SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12396	42764	A	12471	3771	4703	SFQFFCSVFSPSLWFYLLLVFDD
12390	42/04	^A	124/1	3//1	4703	GDVQMRFRCGCPFCLLVFLLTD
						RTLSCRSVGIPCRVRCQCA\LLG
						GASQLGCSGVKGSRDPLEEAV
						CPFS\DL\QLRAGGNTALFKAVR
						QGHLSLQRLLLSF\VCLCPAPRG
						GAYRGGQASLSCGGLHPV*AS\
						RLLCLPKQAWAMAGAPPPASL
						L\PCI\LISDCCASNQRDSVGVGP
						S/ERPGVGYSLVVRRFLSRSEKR
						NIRGG\VTRFSRCRPSPLSLTRK
						GNSLTPCASQSEGNAIGPAFGF
						AHGARTHWALRPTVWHSLCE
						MNPVPQMEMQKSPVFCVAHA
			!			GGCRPEL
12397	42765	Α	12472	1	1038	odeld EE
12398	42766	A	12473	27	428	LADRSAIPGFAIFAVTFLLALVG
12070	1.2700		12 (75	<u> </u>		AVLYLYPWTLLKPC*SHY*GIN
						LVVAV*VKTT*GKNCMKMV*L
	1					IL*RVTLPSS*RFGLRLEKAF*M
						GHLIKT*LGKNNMKMPSCNWS
						LF*GTS*KNEKEGTSVNIFSLTP
12399	42767	A	12474	107	362	NKPCEEFYFLAALIYLLYHGRK
						KADTKIPP*ISRFFLILYPSHELTI
						FIVIHIMCDLGCTWAICFLTTSE
						EVQKKLYEEINQVFG
12400	42768	A	12475	193	637	TSSEVVKKQIAQVHNFAVIMQL
	1					ARENIILSS/EDLLVIKVPLY*GV
	1					NENMLTEVPSFSFFYDVP*NRL
						QLHEGIFILFFPSHVFIK*PI*KAF
						SNLRPNCAMILLEANDFLIIFKC
						TTHYHLCNRLHSKTKHMLAEG
						HVLGLWVGEPFI
12401	42769	Α	12476	1	1930	
12402	42770	Α	12477	1	362	
12403	42771	Α	12478	58	160	GVSLQSTSEEGRGMRQE*TERE
	<u> </u>	<u> </u>				IKVTCRSHYRLA
12404	42772	A	12479	3	410	
12405	42773	A	12480	1	807	
12406	42774	A	12481	154	375	
12407	42775	A	12482	2	512	
12408	42776	C	12483	29	455	
12409	42777	A	12484	209	433	
12410	42778	A	12485	1	1546	
12411	42779	A	12486	3	931 363	
12412	42780	A	12487	1		
12413	42781	A	12488	1	351	
12414	42782 42783	В	12489	82	263 1878	
12415 12416	42784	A	12490 12491	24	242	
	42785	A	l		3558	
12417	142/83	A	12492	11	2230	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleatide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
İ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		, · · · · · · · · · · · · · · · · · · ·
12418	42786	A	12493	2	478	IDICI KODENKENOOO DAAKEE
12110	142700	`	12493	12	470	HNSLKSDENKENSFSADHVTTA
		l				VEKSKESQVTADDLEEEKAKA
	1					ELIMDDDRTVDPLLSKSQSILIS
		1				TSATASSKKTIEDRNIKNKKSTN
						NRASSASARLMTSEFLKKSSSK
						RRTPSTTTSSHYLGTLKVLDQK
12419	40707	<u> </u>	12404			PFTETEA*EPDRADNIRGSCLFR
12419	42787	A	12494	1	1047	
12420	42788	A	12495	1	1055	
	42789	A	12496	212	611	
12422	42790	Α	12497	2	195	YVMGRNDGGYLMIDSKTAEIK
i						FVKNMNRDSTFIVNKTITAEVL
	<u> </u>	 				AID\RYSGKNCLLISWSYICG
12423	42791	Α	12498	1	429	
12424	42792	Α	12499	1	503	
12425	42793	Α	12500	3	825	GTTKWQTVRRQKREWIKFAAA
		1				CREGEDNSKRNPIAK*SDLQNS
Ī						LLPALY*KH\IRSDCESNQKITY
						RISGVGIDRPPYGVFTINPRTGEI
						NITSVVDREITPLFLIYCRALNS
					1	RGEDLERPLELRVKVMDINDN
						APVFSQSVYTASIEENSDANTL
						VVKLCATDADEENHLNSKIAY
	1					KIVSQEPSGAPMFILNRYTGEV
						CTMSSFLDREQHSMYNLVVRG
						SDRDGAADGLSSECDCRIKVLD
						VNDNFPT/FRENFKQNSSRFSNS
						LLSSFDQWRKL
12426	42794	A	12501	986		KQRLTWTFSPGKSETMMGLFP
						RTTGALAIFVVVILVHGELRIET
						KGQYDEEEMTMQQAKRRQKR
					l l	EWVKFAKPCREGEDNSKRNPIA
					ı	KITSDYQATQKITYRISGVGIDQ
						PPFGIFVVDKNTGDINITAIVDR
		l				
						EETPSFLITCRALNAQGLDVEKP
		- 1				LILTVKILDINDNPPVFSQQIFM
					l l	GEIEENSASNSLVMILNATDAD
						EPNHLNSKIAFKIVSQEPAGTPM
						FLLSRNTGEVRTLTNSLDREQA
12427	42795	В	12502	1	3572	SSYRLVVSGA
					594	
					612	
					532	
					735	
		-			400	
- -734	72000	<i>n</i>	14301	J	400	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12433	42801	Α	12508	2	1260	GQAASPASPASPAMAWAA
						LLGLLVALLLLLLSRRRTRRP
						GEPPLDLGSIPWLGYALDFGKD
						AASFLTRMKEKHGDIFTILVGG
						RYVTVLLDPHSYDAVVWEPRT
						RLDFHAYAIFLMERIFDVQLPH
						YSPSDEKARMKLTLLHRELQAL
						TEAMYTNLHAVLLGDATEAGS
						GWHEMGLLDF/CLQLPAQSRLP
						DSLRN*GAATHP*KPGPGPRPLS
}						*CLPHLSPARPAAPQTGPWLPV
						SGGQGPHVQCQKSPVEAAIPSQ
						AGQAGPPEQMAGELPAAPGGD
						GCVRGDAGTGPGAAAVGHTGE
		ŀ				YGSRCLLAPALPSQES*SPGCCP
1						RRAREYPLASGAACLADDHSPT
						EGSRQHTC/DLIAC*VRASGLQL
		Ì				PPSSPARL\GGPGHAHGRRERIQ
						PATW*PPPPLPLPEPPERPRNLH
						RPRGI
12434	42802	Α	12509	3	466	
12435	42803	A	12510	1	1134	
12436	42804	A	12511	2	1211	
12437	42805	Α	12512	2	405	HEMLLASEIKHLPRRLLIGQERE
						TQTSELKIKRRGNEEAPS/PPPSS
						AYERGTKRPDDRYDTPTSKKK
						VRIKDRNKLSTEERRKLFEQEV
			!			AQREAQKQQQQMQNLGMTSP
						LPYDSLGYNAPHHPFAGYPPGY
						PMQA
12438	42806	A	12513	191	898	
12439	42807	A	12514	1	270	
12440	42808	Α	12515	2	1328	
12441	42809	A.	12516	1	2868	
12442	42810	A	12517	1	1254	
12443	42811	A	12518	1	576	

SEQ ID	SEQ ID NO:		-			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	1	07/340,217	sequence	or peptide sequence	deterion, (-possible indeteoride insertion)
12444	42812	Α	12519	1919	3044	HQGPSTPPSWAMSGPPTPLSRE
				ŀ		DWHQGPSTPPSWAMSEPPT/SSI
						QGLASGAVHTILLGDVRATYTS
						IQGVTSGVSQVSRAAQMAVPSS
						RILQLSKPKAPATLLE\EWDPVP
						KPKPHVSDHNRLLHLAKVPRK
						EGSGKKVGAFPEIKGPEAFRDK
						ARAMESQSNDMPFDELLALYG
						YEASDPISDRESEGGDVDPNLP
						DMTLDKEQIAKDLLSGEEEEET
						QSSADDLTPSVTSHEASDLFPN
						RSGCLLAGEAESSRGLLPRAQP
						VPRGAGLADNSRGALLRAHGT
						VRVGTTATVKPADAPPESPRDR
						RSRNDSHRPTGPSESERQPQSN
						QPTLLLRGHGTIRVRTTATVKP
						ADAPAESPRDRRSRNDSHGQSS
12445	42813	Α	12520	413	1412	
12446	42814	A	12521	293	412	NHLSVAGNGSSPCGL*AGALGL
		1				LLWLLASSGHRVLSGMI
12447	42815	Α	12522	2	181	
12448	42816	A	12523	1	277	
12449	42817	Α	12524	1	739	
12450	42818	Α	12525	1	326	
12451	42819	Α	12526	3	566	
12452	42820	Α	12527	1	465	
12453	42821	A_	12528	1	1167	
12454	42822	Α	12529	260	395	EVATATPTLSNHYPDQSAAIN/D
						QG*PLHQQKDHILQKAQMNISI
		<u> </u>				F
12455	42823	A	12530	209	958	FRCVLISFWSHKLHLWYHEGRI
		ŀ				VPSDRIVPPHGIVSSHGVALPQR
						ILMRQFTLLEGLNKDGRFLIQLS
						GTAPWSSYFNSVAKFGVIHRIH
		İ				SFFLSHNGHKAVAPLDLLSNVG
ļ		1				GSCLCGDASSF\TTPSRITMTSK
1			1			SWLARSVAEMSYR/WTALVNH
						VADDQRGHQKNVIALDWVINR
						NVQLVNRNHLPFRCSCFLDHLD
						AHRGTNDHAFPQVTHSEHEAQ
						LAISDGNHSMAAENQSFCPSVS
10:55	10001	ļ.	10555	ļ	1065	LSCFHENAA
12456	42824	A	12531	1	1965	
12457	42825	C	12532	520	684	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12458	42826	A	12533	701	1307	GRHLTTNHTAAVVVFPFRPLCS SFFRYSGFRGVAGATASGRGQG PAG*DNCQRAWRGAFYKFQAT LSAGAGTRAEEGSDGDGEAAE PGAGTAREPARLRGYRSRRFVL DGRRCCLCYFKNPHDPLPLGHL DIAEACFSYQRADEAAEQPAHF QVRSAGAATVLEVGTRALRGV HLLRAHRGIWAQAWSPVSFLQ AVLPAWLL
12459	42827	Α	12534	1	1356	
12460	42828	В	12535	1	1449	
12461	42829	Α	12536	3	2449	
12462	42830	Α	12537	1	3327	
12463	42831	Α	12538	1	1359	
12464	42832	Α	12539	1	672	
12465	42833	Α	12540	1	1281	
12466	42834	Α	12541	1	399	
12467	42835	Α	12542	1	399	
12468	42836	A	12543	171	426	PVSCATGEEEDNVLFPKGDDLP R**PGKLWCEFHASSISQALGQ HPRRSVLDTDQSHVPLVRRKTT CFSQRAMISPDDDPSQEAGALS RNNLLAPYSALTFMENGNCLL QLFQLGKLLVQASHLHGQLLV FVQKIIISM
12469	42837	Α	12544	1	988	
12470	42838	A	12545	3	129	LFHPCQDSQQHH*CVCCRLTGH GAA*VHGPCQAVQTYRASH
12471	42839	С	12546	532	1101	
12472	42840	A	12547	197	355	
12473	42841	Α	12548	1	1275	
12474	42842	Α	12549	3	131	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12475	42843	I _A	12550	1746	3504	ESDPSNSTTSMDRTLL/MSSIVP
12473	172075	ľ``	12330	1740	3301	RLPSACRILPTGFTIPE*SCVHRT
]				ASLIPPLPPGSRKYSPPLPLNSLI
						FLKN/SAYTAVPALQTDWATSP
		İ				ISLHLRTSFNSPHLYPPEELIYFL
		l				DRSSKTSPDISHQQAAALLRTY
		İ				LKNLSPYINSTPPIFGPLTTQTTI
		l				PVAAPLCISWQRPTGIPLGNLSP
		İ				SRCSFTLHLRSPTTNINETIGAF
						QLHITDKPSINTDKLKNISSNYC
						LGRHLPCISLHPWLSSPCSSDSP
						PRPSSCLLIPSPENNSERLLVDT
						l l
						RRFLIHHENRTFPSTQLPHQSPL QPLTAAALAGSLGVWVQDTPF
						1
						STPSHLFTLHLQFCLAQGLFFLC
						GSSTYMCLPANWTGTCTLVFL
		l				TPKIQFANGTEELPVPLMTPTQ
						QKRVIPLIPLMVGLGLSASTVA
						LGTGIAGISTSVMTFRSLSNDFS
						ASITDISQTLSVLQAQVDSLAA
						VVLQGQLRGLDLLTAEKGGLCI
						FLNEECCFYLNQSGLVYDNIKK
		1				LKDRAQKLANQASNYAESPWA
						LSNWMSWVLPIVSPLIPIFLLLL
						FGPCIFRLVSQFIQNRIQAITNHS
	12271	<u> </u>				IRQMFLLTSPQYHPLPQDLPSA
12476	42844	A	12551	2	394	
12477	42845	A	12552	1	882	
12478	42846	A	12553	45	409	
12479	42847	A	12554	1	1035	
12480	42848	A	12555	118	312	
12481	42849	A	12556	1	1083	
12482	42850	В	12557	1	1122	
12483	42851	Α	12558	97	439	
12484	42852	A	12559	1	450	
12485	42853	Α	12560	1	471	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12486	42854	Α	12561	2	1407	WGKNRAEKLETLKSRVPLLLQ RNAVPHQQWNKAGRRMTLTS* EKKASDD/PNYSKLQEEIQTKG KEVKHFEKNLDECITRITNREK CLKELMELKAKARELREECRSL RSQCDQLEERVSVMEEIQTTIRE YYKHLYENKLQNLEEMDKFLD TYTLRRLNQEEVESLNRPITGSE IVAIINSLPTKKSPGPDGFTAKF YQRYKEELIPFLLKLFQSIEKEGI LPNSFYEASIILIPKQGRDTTKK ENFRPIFLMNINAKILNKILANRI QQHIKKLIHHDQVGFIPGMQG WFNICKSRNVIHHINRTKDKNH MIISIDAEKAFDKIQQPFMLKTL NKLGIDETYLKIIRAIYDKPTAN IILNGQKLEAFPLKTGTRQGCPL SPLLFNIVLEVLARAIRQEKEIK GIQLGKEEVKLSLFADDMIVYL
						ENPIVSAQNLFELISNFSKVSGY KINVQISQAFLYTNYRQSAKS
12487	42855	Α	12562	1	846	
12488	42856	Α	12563	1	1281	
12489	42857	В	12564	1	274	
12490	42858	В	12565	1	2265	
12491	42859	Α	12566	1	2742	
12492	42860	Α	12567	1	1272	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12493	42861	Α	12568	222	2130	RKKNKKK*TKPPRNMGLREKT
						KSMSDWCT*K*RGEWNQV/AK
		<u>.</u>				TLCRILSRRT/YPNLARQADIQIQ
						EIQRTPQRYSSRRATPKHIIVRF
		İ	,			TKVEMKEKMLRAAREKGRVT
						HKGKPIRLTADLSAETLQARRE
		•				WGQIFNILKEKNFQPRISYPAKL
						SFISEGEIKPFTDKQMLRDFVTT
						RPALKELLKEVLNMERNNRKK
						LNRPITGSEIVAIINSLPTKKSPG
		·				PDGFTAEFYQRYKEELVPFLLK
						LFQSKEKEGIHPNSFDEASIILIP
						KAGRDTTTKDNFRPISLMNIDA
						KILNKILANQIQQHIKKLIHHDQ
						VGFIPGMQGWFNICQSINVIQHI
						NRTKDKNHVIISIDAEKAFDKIQ
						QRFMLKTLNKLGIDRMYLKIIR
1						AIYDKPTANIILNGQKLEAFPLK
]					MGTRQGCPLSPLLFNIVLEVLA
						RAIRQEKEIKHIQLGKKEVILSL
						FADDMIVYLENSIVSAQNLLKLI
						SNFSKVSGYKINVQKSHAFLHT
		•	1			NNRQTESQIMSELPFTIASKRIK
						YLGIQLTKDVKDLFKENYKPLL
						NEIKEDTKKWKHIPCSWVGRIN
						IVKMAILPKVIYRFNAIPIKLPM
						TFFTELEKTTLKFIWNQKRAHT
						AKSILSQKNKAGGITLPDFQLY
12404	42862	Δ.	12569	1	474	CKATVTKTAWY
12494 12495	42863	A A	12570	1	1197	· -
12496	42864	A	12571	1	1518	
12497	42865	A	12572	1	3189	
12498	42866	A	12573	1	2199	
12499	42867	A	12574	3	2820	ENKDTTYQNL\WDAFKAVCRG
12477	12007	, ,	12374	ľ	2020	KFIALNAHKRKQERSKIDTLTS
						QLKELEKQEQTHSKASRRQEIT
						KIRAELKEIETQKSLQKINESRS
						WFFERINNIDRPLARLIKKKREK
						NQIDTIKNDKGDITTDPTEIQTTI
ŀ						
						REYYKHLYANKLKNLEEMDKF
						LDTYTLPRLNQEEVESLNRPITG
						SEIVTIINSLPTKKSPGPDGFTAE
						FYQRYKEELVPFLLKLFQSIEKE
						GILPNSFYEASIILIPKPGRDTTK
	12.2.65	<u> </u>				KENFR
12500	42868	В	12575	1	1593	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12501	42869	Α	12576	1	3549	MGDFNTPLSTLDRSRRQKVNK
						DTQELNSALHQVDLIDIYRTLH
i						HKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTONRSTTWKLN
						NLLLNDYWIHNKMKAEIKMFF
1						ETNKNKDTTYQNLWDAFKAV
						CRGKFIALNAHKRKQERSKIDT
1						LTSQLKELEKQEQTHSKASRRQ
						EVTKIRAELKEIETQKTLQKINE
ł						SRSWFFERINKIDRMLARLIKK
						KREKNQIDTIKNDK
12502	42870	Α	12577	1	2724	
12503	42871	A	12578	1	3567	
12504	42872	В	12579	709	2064	
12505	42873	A	12580	1	3370	
12506	42874	Α	12581	1	2024	MGFYHVGQAGTELLTSNDPLT
						SASQSARITGKALQKPTEVPVP
						YEKMLQDQSALIVQGLPEGVA
						FKHPENYDLATLKWILENTAGI
						SFIINSATIEPPAAQPMSFTHMG
						HSLCVLVVETHAKPVAVQSSSF
1						TRNPEVISLEVAAVTVKEESED
ł						PDYDYITFKDLPGIAYETKAYV
					1	LWPQLSKICSKNGDKNKAYRG
					ł	QQCRLRQQNPLAALTGKISVSV
						YVSHPSLRSVSIGWTRQVARQT
						MQKDRDGPLKNEGEKNCAGSE
					İ	KNTLTSQLKELEKQEQTHSKAS
						RRQEITKIRAELKEIETQKTVEK
						INESRSWFFERINKIDRPLARLIK
						KKREKNQVDAIKNDKGDITTDP
						TEIQTTIREYYKHLYANKLENV
						EEMDKFLDTYTLPRLNQEEVES
						LNRPITGSEIVAIINSLPTKKSPG
						PDGFTAEFYQRYKEELVPFLLK
						LFQSIEKEGILPNSFYEASIILIAK
						PGRDTTKKENFRPISLMNIDAKI
						LNKILAKRIQQHIKKLIHHDQV
	!					GFIPGMQGWFNIRKSINVIQHIN
						RAKDKKHMIISIDAEKAFDKIQ
						QPFMLKSLNKL/DIGKNYFKVH MGPKKSPCRQVNPKPKEQSWR
						1
						HHST*LQTILQGYSNQNSMVLV
						PKQRYRSMEQNTALRNNATYL QLSDL*QT*EKQAMGKGFPI**
12507	42875	Α	12582	722	4214	Arania di Evaniara
12307	720/3	<u> </u>	12302	144	7617	

1 1	of peptide	hod	' TICCT'			
s	` '	nou	in USSN	location of first	codon for last amino acid	
: 1	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12508 4	42876	A	12583	228	1152	KNRDYNKLSLRPQCNQTRTQN
						*ESHSKPLNYMETEQPAPE*LL
						DT*R\MKAEIKMFFETNENKDT
						TYQNLWDAFKATASKTNKEKE
						KNQIDTIKNDKGDITTDPTEIQT
						TIREYYKHLYANKLENLEEMD
						KFLDTYTLPRLNQEEVESLNRPI
						TGAEIVAIINSLPTKKSPGPDGF
						TAEFYQRYKEELIGWLRESVSK
						ICLKFDHQVQVITGRLIIRFQIAF
						GTVVLCQLTRHIDVFVGKRLA
						NARFAPSLGGAPDLFTLKPCFV
						AVETQGDLPQATTVVDIDGCL
						GKPANVQVALDLDVKGFQQW
12509 4	42877	A	12584	1	2028	
	42878	Α	12585	2	2264	
	42879	Α	12586	1	2130	
12512 4	42880	Α	12587	1	2900	MENDFDELREEGFRQSNYSELR
						EDIQTKGKEVENFEKNLEECITR
						ITNTEKCLKELMELKTKAQELR
						EEWRSLRSRCDQLEERVSAME
						DEMNEMKGEGKFREKRIKRNE
						QSLQEIWDYVKRPNLCLIGVPE
						RDGQNGTKLENTLQDVIQENFP
						NLARQANVQIQEIQRTPQRYSS
						RRATPRHIIVRFTKVEMKEKML
						RAAREKEIQTTISEYYKHLYTN
						KLENLEEMDKFLDTYTLPTLNQ
12512	42001		12500	1	2745	EEVESLNRPITGAE
	42881	A	12588	1	2745	
	42882 42883	B A	12589 12590	1090	3288 2526	
	42884	A	12591	1090	3242	
	42885	A	12591	3862	8976	RAKSPANIIMTGSNSHITILTLN
12317	42003	^	12392	3602	6970	VNGLNSPIKRHRLASWIKSQDP
						SVCCIQETHLMCRDTHRLKIKG
						WRKIYQANGKQKKAGVAILVS
						DKTDFKPTKIKRDKEGHYMMV
						KGSIQQEELTILNMYAPNTGAP
						RFIKQVLSDLQRDLDSHTLIMG
						DFNTPLSTLDRSTRQKVNKDTQ
	1					ELNSALHQADLIDIYRTLHPKST
			,			EYTFFSAPHHSYSKIDHILGSEA
						LLSKCKRTEIITNYLSDHSAIKL
	İ					ELRIKNLTQSR
12518 4	42886	A	12593	1	1000	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12519	42887	A	12594	1	979	MNNAKENFLGRFQDGRIGTAP
						VYSPQHQRRRRRVISALPTEPPL
						VIPRQTGFGVDLQQTPTDLQLR
						VLTVRRKTTKQEGHSTKTPSVR
						YHHQRPKEDKTTKMGRNQSRK
		1				AENSKNESASSPPKECSSSPATE
						QSWMENDFDKYTEVGFRQLVI
						TNFSELKEDVQTHHKEAKNLE
						KRLDEWLTRINSIENTLIDLMEL
						KTMARELRDSCTSFSRQFDQVE
						ERVSVIEDQMNEMKREEKFRE
						KKMLEVLPRAIRQEKEIKGIQL
						GKEEVKLSLFADNMTVYLENPI
						IS\PKISLS**ATSAKSQDIKSMC
		<u> </u>				KNHKHSYTPITDKQRAKS
12520	42888	Α	12595	1140	2210	KPWHKNFVTHAQASIADLIKW
						KKGYQ*LKIKLMK*SKKTRNTN
						YHQNTINISMQIN*KI*KKWRNS
						WTHTLSQY*TRKKLNL*IDQ*Q
						ALKL/QAIIYSLPTKNVQGQMD
						SQSNTTRVWGILARAIRQEKEI
						KDIQLENEEVKLSLFADDMTVY
						LENSIVSAQNLLKLRIKYLGIQL
						TTDVKDLFKENYKPLLNKIKED
						TNKWKNIPCSWIRGINIMKMAI
				-		LPKLIYRFNAIPIKLPMTFFTELE
						KATLKFIWNQKRAHIAKTILSK
						KNKAGGIMLPDFKLYYKATAT
]		KAAWYWYQNRDIDKKLTQIYK KKTNNPIKKWAKDMNRHLSKE
						i i
						DIYGANRHMKKCSSSLVIREKQ IKTTMR
12521	42889	A	12596	520	697	GLCSVPLLCISVLVPVPCCFGYC
12321	42009	^	12370	320	057	SLVV*FEVR*SDASSFVLLA*DF
						LGNAVSFLVPYEL
12522	42890	A	12597	1	1410	DOINTYOLDYLIDD
12523	42891	A	12598	1	1461	
12524	42892	A	12599	77	457	QALKLRQ*LIAYQPKRVQDQM
_						DSQTNSARGTKRSWYHSF*NYS
			1			N/PTEKEGILLNSFYEAGIILIPKP
		l				GRDTTKKENFRPISLMSINAKIL
						NKILANRILQHIKKLI\ITIKSASS
						LGCKAGSTYANQ
L		<u> </u>	L	<u> </u>	1	

DE CODON, /=possible nucleotide ion, \=possible nucleotide insertion) DENTPLSTLDRSTRQKVNK ELNSALHQADLIDIYRTLHP EYTFFSAPHHTYSKIDHILG ALLSKCKRTEIITNCLSDHSA ELRIKKLTQNCSTTWKLNN
DFNTPLSTLDRSTRQKVNK ELNSALHQADLIDIYRTLHP EYTFFSAPHHTYSKIDHILG ALLSKCKRTEIITNCLSDHSA
ELNSALHQADLIDIYRTLHP EYTFFSAPHHTYSKIDHILG ALLSKCKRTEIITNCLSDHSA
ELNSALHQADLIDIYRTLHP EYTFFSAPHHTYSKIDHILG LLSKCKRTEIITNCLSDHSA
ELNSALHQADLIDIYRTLHP EYTFFSAPHHTYSKIDHILG LLSKCKRTEIITNCLSDHSA
EYTFFSAPHHTYSKIDHILG LLSKCKRTEIITNCLSDHSA
LLSKCKRTEIITNCLSDHSA
CERTIFICE I WILDING
NDYWEHNKMKAEIKMFF
IKNKDTTYQNLWDTLKAVC
KFKALNAHKRKKERSQIDTL
-
LKELEKQEQTHSKASRRQE
IRAELKETETQKTLQRLMN
VLEVLARAIRQEKEIKGIQL
EEVKLSLFADDMIVYLENP\
QPKISLS**ATSAKSQDIKSIY
HQHSYTPIRDKQRAKS
KFLDTYTLPRLNQEEVESLN
rgseieaiinslptqkssgpd
'AEFYQRYKEELVPFLLKLF
EKEGILPNSFYEASIILIPKPG
THKKNNFRPISLMNIDAKIL
LANRIQQHIKKLIHQDQVGI
MQSWFNIHKSINVIQHINRT
(NHMIISIDAEKAFDKIQQPF
KTLNKLENKIPKNPTYKGCE
QGELQITAQRNKGRHKQM
IPMLMDRKNEYHKNGHTPK
RFNVIPIKLPMTFFSELEKST
IWNQKRARIAKTILSQKNK
GIMLPDFKLYYKATVTKTA
WYQNRDIDQWNRTEPSEM
IYNHLIFDKPDKNKQWGK
FNKWCWENWLAIGRQLKL
LTPYTKINSRWIKDLNVRP
KTLEENLGNTIQDISMGKDF
CTPKAMATKAKMDKWDLI
SFCTAKETTIROSWPEMTK
SSNTDVKNWKHGKNNIGR
SVNTCKMDSGLVLERWM
NPGFPLSIDAKCHKDLPRDI
SEKGVDFVLNYSKANMR
GSMFL*FIESF*HEWLLNFV
FCIY*DNHVVFVFGSVYML
71111
LSQPKISLS**ATTAKSQDT
YKNQQHSYT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12530	42898	A	12605	77	1866	QAPKLRQ*LIAYQPKKVQDQTD
12330	42070	``	12003	1''	1000	SQPNSTRGTKKSWYHSF*NFSN/
						PIEKEGIFPNLFYEANIILIPKPGR
						DTTKKENFRPISLMNVNAKILN
						KLLANQIQQHIKKLIHHNQVGFI
1						PGMQGWFNICKSINVIQHINRT
						KDKNHRSISIDAEKAFNKIQQPF
1						MLKTLNKLDDMIVYLENPIVSA
						KNLLKLISNFSKVSGYKINVQK
						1
						SQAFLYTNNRQAESQIMSAVSF
						TIASKRIKYLGIQLTRDVKDLFK
						ENYKPVLNKIKEDTNKWKNIPS
]		LWIGRVNIMKMAILPKVIYRFN
				1		AIPIKLPMTFFTELEKTTLKFIW
}			:			NQKRAHIAKTILSKKNKAGGIT
						LPDFKLYYKATVTKTAWHWY
						QNRDIDQWNRTEPSEIIPHIYNH
						LIFDKLDKNKTLGKDYLFNKRC
						WENWLAICRKLKLDPFLTRYT
				1		KINSRWIKDLNIRPKTIKTLEEN
						LGDTIQDIGMGKDFITKTPKAM
						ATKAKIDKWDLIKLKSFCTAKE
						TTIRVNRQPTEWENIFAIYPSDK
						GLISRIYKELKQIYKKKSNNPIK
						KWAKDMNRYFSKEDIYAANR
						HMKKCSSSLALREMQIKTTMR
						YHLIPVRMAIIKKSGNNRCW
12531	42899	Α	12606	67	243	GLCCVPLIYISVLVPVPCCFGYC
]		SLVV*FEVR*HDASSFVLLA*D*
						LGDVGSFLVPYEL
12532	42900	Α	12607	1554	1692	
12533	42901	A	12608	640	816	GLCSVALVCISVLVPVPCCFGY
						CSLVV*FEVR*HDASSFVLFA*D
			<u> </u>			*LGNVDSLLVPYEL
12534	42902	Α	12609	1626	1739	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12535	42903	A	12610	1	2014	MKREGSLEKKRIKRNEQSLQEI
						CDYVKRANLRLIGGPESDGENG
						TKLENTLQDMQEKFPNLARQA
		ŀ				NIQIQEIQRTPQRYSSRRATPRHI
						IVRFTKVEMKEKMLRAAREKG
		ŀ				QVTHKGKSIRLTADLSAETLQA
:						RREWGPIFNILKEKNFQPRISYP
		l				GKLSFISEGEIKSFTDKQMLRDF
						VTTRPALKELLKEALNMEKNK
						QYQPPQKHAKLTREARANTFK
						PSRRREITKIRAELKEIETPKTAQ
						KINESRSWFSERINKIDRPLARL
						RKKKREKNQIDTIKNDEGDITT
						HPTEIHTIIREYYKHLYANKLEN
				ŀ		LEEMDKFLDTYTLPRPNQEEVE
				}		SLNGPIAGSEIQAIINSLPTKKSP
						GPDGFYQRYKEELVPFRLKLFQ
		1				SIEKEGILPNSFYEASIILIPKPGR
						DTTKKENFRPISLLNINAKILNKI
						LANRIQQHIKKLMHHDQVGFIP
						GMQGWFNIRKSINVIQHINRTK
						DKNHMIISIDAEKAFDKIQQPF
				ŀ		MLKTLNKLGIDGTYHKIIRANY
				ŀ		DKPTANIILNGQKLEAFPLKTGT
				ŀ		RQGCPLSPLLFNIVLEVLARAIR
						QEKETEGIQLGKEEVKLSLFAD
						DMIVYLENP\MSQPKISLS**ATS
						AKSQDTKSMCKNHKYSYTPIT
						DKQRAKSSVNSPFTIASKRIKYL
						GIQLTRYVKDLFKERSMK
12536	42904	Α	12611	1	700	MDKFLDTYTLPRLNQEEVESLN
						RPITGSEIEAIINSLSTKKSPGPD
						GFIAQFYQRTKDKNHMIISIDAE
						KAFDKIQQHFMLKTLNKLGIDG
						TYLKIIRAVYDKPTASIILNGQK
						LEAFPLKTGTRQGCPLSPLLFNI
				}		VLEVLARAIRQEKEIKGIQLGKE
						EVKLSLFADDMIVYLENP\MSQ
						PKISLS**ANSAKSQDIKSMCKN
						LKHSYTLITDKQRAKSSVNSHS
			L	<u>L.</u> .		QLLQRE
12537	42905	Α	12613	1	1765	
12538	42906	Α	12614	1	2310	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12539	42907	A	12615		836	MENEFDELTEVHFRRWVITNSS ELKEHVLTQCEEAKNLARRNK LENLEEMDKFLATYTFPSLNQE EVESLNRPITSSEIEAAINSLPTK KHPGPDGFTAKFYQRYKEELHI NRAKDKNHMIISIDAEKAFDKI QHPVKLKILNKLGIDGTYLKIIR AIYDKPTANLILNGQKLEAFPL KTGTRQGCPLSPLLFNIVLEVLA RAIRQEKEIKGIQLGKEEVKLSL FADD\RLYI*GTPLSQPKISLS** ATSAKSQDTKSMYKNHKHSYT PTTDKQRAKS
12540	42908	A	12616	355	4502	QSARFRGRRTLRRPGVGYSLVV *FEVR*CDASSFVLLA*DCLGN AGSFLVPYEL
12541	42909	Α	12617	248	362	
12542	42910	A	12618	529	680	YYTRCSKWWNCSVLP*IYIYNF HLPHGKLKNTDYIKKKLHDFFF KNSKHI
12543	42911	A	12619	1	641	
12544	42912	В	12620	36	48	
12545	42913	Α	12621	1	326	
12546	42914	Α	12622	I	444	PGSTHASAHACERLQTRTLRAF SSQRRFPATASRASLSSNMAKIS SPTETERCIESLIAVFQKYAGKD GYNYTLSKTEFLSFMNTELAAF TKNQKDPGCP*PHDGRKLGQPT VDGQLDFSEFLNLIGG\QAMAC HDSFLKAVPSQKRT
12547	42915	A	12623	2	6008	
12548	42916	A	12624	530	856	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12549	42917	A	12625		952	MAARGTGDKVKPGCEEAGAAF ELPPSVYAAALHESHQHQPCPA GGARAQRVPGGIVTEPLTCTPP KDMVFLSDDGDDDDDDDDDDDDDDDDDDDDDDDDDDDDDDDDDD
12550	42918	A	12626	3	380	
12551	42919	Α	12627	5	765	RQWEVGRWSGTTIFSCFCKKTL YFTNPSPSSSSLVTVILRREVTH/ CGGAPEQRYQIIPVCVAARLPT RAQDVLDAHLSEVNAVRFGPN SSLLATGGADRLIHLWNVVGSR LEANQTLEGAGGSITSVDFDPS VRNSAPVAWDCALSDLHTGQG YQVLAATYNQAAQLWKVGEA QSKETLSGHKDKVTAAKFKLT RHQAVTGSRDRTVKEWDLGRA YSPKRQRPSRTGHDDDGGFVE KKRGKCGEKRDQIVTVSV
12552	42920	В	12628	1	2802	
12553	42921	A	12629	458	708	QNRHRHHGPFSMVAVSSQLLD TPAERLSLHTWKIAQRPGRGTP HFPDGAV/VQAETLPQTGRPGR GTPHFPDGAASSFGYCSLVA
12554	42922	A	12630	2300	6491	EGLEALIHQERLSYWANQVSED RPVCKAIIQGKQFEGLVDTGAD VSIIALNQWPKNWPKQKAVTG LVGIGTASEVYQSMEILHCLGP DNQESTVQPMITSIPLNLWGRD LLQQWGAEITMPAPLYSPTSQK IMTKMGYIPGKGLG\KNEDGIK VPVEAKINQEREGI\GYPFLGAV TVEPPKPIPLTWKTEKPVWVNQ WPLPKQKLEALHLLANEQLEK GHIEPSFSPWNSPVFVIQKKSGK WHTLTDLRAVNAV

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
12555	142022		12621	1	434	MTYCKNIK CUTIK CCK K CTEKK
12555	42923	A	12631	1	434	MTVSKNKCHTKGGKKGTEKK
	1					VVASFSKIYWYNVKAPAMFSIR
						NIGKALVIGTHGTDFVSDGFKG
ļ						GLIEIPDSIGKDKEKACRSIYPLH
			'			DVFVRKVKMLKKPKFELGKLM
İ						ELHGEGRSSGKATGNESGA/KV
						E*ADGYGSPAQKSF
12556	42924	Α	12632	1	642	
12557	42925	Α	12633	260	534	
12558	42926	A	12634	1	438	
12559	42927	С	12635	391	750	
12560	42928	A	12636	199	451	
12561	42929	Α	12637	120	992	RDRIPGPVCKGKWKNKERILIFS
]	ŀ				SRGINFRTRHLMQDLRMLMPH
		1				SK\HDTNMARSDNLSVINEGCE
]					MNNWLRCIYFEAKKKQDLYM
		1				WLSNSPDGPSAKFLVQNIHTLA
		į				ELKMTGNCLKGSRPLLSFDPAF
						DELPHYALLKELLIQIFSTPRYH
		ļ				PKSQPFVDHVFTFTILDNRIWFR
		İ				NFQIIEEDAALVEIGPRFVLNLIK
						IFQGSFGGPTLYENPHYQSPNM
						HRRVIRSITAAKYREKQQVKDV
						QKLRKKEPKTLLPHDPAADVF
		1				VTPAEEKPIEIQWVKPETKVDL
			Ì			KASV
12562	42930	Α	12638	1	342	
12563	42931	Α	12639	1	580	PGWEKRMSRSS/GVNTQEALPT
						AAIPTDAKVRVYYFNHITNASQ
						WERPIGNSSSGGKNGQGEPARV
	İ					RCSHLLVKHSQSRRPSSWRQEK
						ITRTKEE\ALELINGYIQKIKSGE
						EDFE\SLASQFSDCSSAKA\RGD
						LG\SFSRGQMQKPF*RTPRFAL\
						RTGGDERGPCFTD\SGIHIILPHL
						SEGGKPRPGLGAGQGG
12564	42932	A	12640	3	116	
12565	42933	A	12641	2	253	
12566	42934	A	12642	3	512	EDYLERKRQ/VPTECQTHSKLG
					· -	GPMTLKREVKP/KVDVSPSKKG
						PL\QHHTLL\VCHVRIFYPGSIQ/
		1				VRWFLNGRKKTAGVVSPNLIR
			-			NGDWT/FQILVMLEM/TPQQGD
						VYTCQVEHTSLDSPVTVEWKA
				1		QSDSARSKTLTGAGGFVLGLIIC
						GVGIFMHRRSKK\FNEDLHKQG
12567	42935	A	12643	1	363	O VOIEWITIKKSKK/FNEDLHKQU
12568	42936	В	12644	70	1229	
12569	42937	A	12645	1	430	
12570	42938	B	12646	1	1321	
12571	42939	A	12647	3	446	
123/1	12/3/	1' 1	1.207/	<u></u>	. 70	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12572	42940	Α	12648	1	417	
12573	42941	Α	12649	269	452	TILRSFGYFFRRINIMLTEEKIKS DHRKISQLIFLQKLCQTLQSENV SQGGMLYRAKA*H*T*TLVWV *HVMSYPHKGLGLMLSLCSVK HSTLRDVFTL
12574	42942	Α	12650	83	1385	
12575	42943	A	12651		974	MNPLECVDRSSQPWGNPTWTS SGMRSVTFLVGFKLTFLALAQT LNAAASLTGQELLAGFPPGVINI LPGYGPTAGAAIASHIGIDKIAF TGSTEKEAELHSWIKGLTCISIE PDFQKSLKDGIILCILMNKLQPA SFPKINHSMQKWHQLENLSNFI KAVVSYDMNPMDLFKANNLFE SGNMRHMQVSLFALAGKVKT KRPQSGVDIGIKYSEKQEQNFS DAAMKAGQCVIGLQMGTNKC TSQLGMTVYGMGTNNCASQVS MTAPGTRQHICDTKLGTDKCG NSSMSLQMGYT/QGAANQPRV CEWQRGIRQQGFLPPCLGPSQE
12576	42944	Α	12652	1	1103	MNPLECVDRSSQPWGNPTWTS SGMRSVTFLVGFKLTFLALAQT LNAAASLTGQELLWNFPLLMF AWKIAPALCCGNTVVIKPAEQT PLSALYMGALIKEAGFPPGVINI LPGYGPTAGAAIASHIGIDKIAF TGSTEKEAELHSWIKGLTCISIE PDFQKSLKDGIILCILMNKLQPA SFPKINHSMQKWHQLENLSNFI KAVVSYDMNPMDLFKANNLFE SGNMRHMQVSLFALAGKVKT KRPQSGVDIGIKYSEKQEQNFS DAAMKAGQCVIGLQMGTNKC TSQLGMTVYGMGTNNCASQVS MTAPGTRQHICDTKLGTDKCG NSSMSLQMGYT/QGAANQPRV CEWQRGIRQQGFLPPCLGPSQE
12577	42945	Α	12653	1	729	
12578	42946	Α	12654	259	390	SFCSANFEVFRLTAQQ*FVR*H ALSHIF*G*V*SFPKSTKRVNF
12579	42947	Α	12655	68	735	
12580	42948	Α	12656	1	825	
12581	42949	Α	12657	3	470	
12582	42950	Α	12658	1	1683	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
12583	42951	A	12659	1	1748	MNKPYIRNLSENAAPLDLEGIL
						KDAKIPVSGPFLVKIGYAFVDC
						PDEIWALKAIEALSGKIELHGKP
						IEVEHSVPKRQRIRKLQIRNIPPH
						LQWEVLDS*LVQYGVVESCQQ
						VNTDSETAVVNVTYSSKDQAR
						QALDK/LLDKLNGFQLENFTLK
						VAYIPDEMATQQNPFLQPRGRR
						GLGQRGSSRQGSPGSVSKQKPC
						DLPLCLLVPTQFVGAIIGKEGAT
		1				IRNITKQTQSKIDVHRKENAGA
						AEKSITILSTPEGTSAACKSILEI
						MRKETQDVKFTEEIPLKILAQN
ļ						NFVGRLIGKEGRNLKKIEQDTD
						TKITISPLQELMQYNPEHTITAK
						GNVETCAKAEEEIMKKIRER*E
						NDITSM/N/LQAHLIPGLNLNAL
			1			GLFPPTSGMPPPTSGPPSAMTPP
		1				SPQSEQSESETVHLFIPALSVRA
						LISKQGQHIKQLSRFAGASSKIA
						PVEAPDAKVRMVMIAGSPEAR
						FKAQGRIYGKIKEENFVSPKEE
		1				VKLEAHIRVPSFAAGRVIGKGG
						KTVNELQNFSSAEVVVPRDQTP
						DENDQVVVKITGHFYACQVAQ
10504	10050		12660		002	RKIQEILTQVKQHQQQKALQSG
12584	42952 42953	A B	12660 12661	13	803 330	
12586	42954	C	12662	127	342	
12587	42955	A	12663	324	491	
12588	42956	A	12664	1	891	-
12589	42957	В	12665	1	1869	
12590	42958	A	12666	2	136	
12591	42959	A	12667	108	355	-
12592	42960	Α	12668	2	811	FGMRGSRGGWAGEMAASGES
						GTSGGGGSTEEAFMTFYSEVKQ
						IEKRDSVLTSKNQIERLTRPGSS
						YFNLNPFEVLQIDPEVTDEEIKK
						RFRQLSILVHPDKNQDDADRA\
						QKAFEAVDKAYKLLLDQEQKK
						RALDVIQAGKEYVEHTVKERK
						KQLKKEGKPTIVEEDDPELFKQ
					İ	AVYKQTMKLFAELEIKRKERE
						AKEMHERKRQREEEIEAQEKA
						KR/EEREWQKNFEES\RDGRVD
						SWRNF\QAQFRRGKKEKKNRTF
		L				LRPPESKNGSNVE
12593	42961	Α	12669	731	850	
12594	42962	Α	12670	201	377	LMTLPCKIPSPGSSWLKISPIEHL
						MTPTPASQRTTPFDCNFPLPTQI
	<u> </u>	<u> </u>				L*NGPTPISLH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
12595	42963	A	12671	742	2982	KIFPFDASFLPCLHKFNLKGKG
						NSKGFKMAEKFESLMNIHGFDL
		:				GSRYMDLKPLGCGGNGLVFSA
						VDNDCDKRVAIKKIVLTDPQSV
						KHALREIKIIRRLDHDNIVKVFE
						ILGPSGSQLTDDVGSLTELNSV
						YIVQEYMETDLANVLEQGPLLE
	ŀ					EHARLFMYQLLRGLKYIHSAN
						VLHRDLKPANLFINTEDLVLKI
						GDFGLARIMDPHYSHKGHLSE
						GLVTKWYRSPRLLLSPNNYTK
						AIDMWAAGCIFAEMLTGKTLF
		į				AGAHELEQMQLILESIPVVHEE
						DRQELLSVIPVYIRNDMTEPHK
						PLTQLLPGISREALDFLEQILTFS
						PMDRLTAEEALSHPYMSIYSFP
						MDEPISSHPFHIEDEVDDILLMD
						ETHSHIYNWERYHDCQFSEHD
						WPVHNNFDIDEVQLDPRALSD
į						VTDEEEVQVDPRKYLDGDREK
						YLEDPAFDTNYSTEPCWQYSD
			ŀ			HHENKYCDLECSHTCNYKTRS
				-		SSYLDNLVWRESEVNHYYEPK
						LIIDLSNWKEQSKEKSDKKGKS
						KCERNGLVKAQIALEEASQQLA
						GKEREKNQGFDFDSFIAGTIHL\
						SSQHEPTDVG\DKLNDLNSSVS
						QLELKSLISKSVSQEKQEKGMA
						NLAQLEALYQSSWDSQFVSGG
		İ				EDCFFINQFCEVRKDEQVEKEN
		-				TYTSYLDKFFSRKEDTEMLET\E
						PVEDGKLGERGHEEGFLNNSGE
12596	42964	Ā	12672	826	1041	WKITVKGGVFLWWAGVGDTK
		1				VLSGGAF*ARMSQEEDFHKVM
						SSVKARTGHLHFFCGGMSSVK
						AGQGPFSVLL
12597	42965	A	12673	1	1482	
12598	42966	Α	12674	3	488	
12599	42967	Α	12675	2	114	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12600	42968	A	12676		897	MSNRVVCREASHAGSWYTASG PQLNAQLEGWLSQVQSTKRPA RAIIAPHAGYTYCGSCAAHAYK QVDPSITRRIFILGPSHHVPLSRC ALSSVDIYRTPLYDLRIDQKIYG ELWKTGMFERMSLQTDEDEHS IEMHLPYTAKAMESHKDEFTIIP VLVGALSESKEQEFGKLFSKYL ADPSNLFVVSSDFCHWGQRFR Y\SYYDESQGEIY*SIEHLDKMG MSIIEQLDPVSFSNYLKKYHNTI CGRHPIG\VLLNAITELQKNGM NMSFS\FLNYAQSSQC\RNWQD SSVSYAAGALTVH
12601	42969	Α	12677	2	114	
12602	42970	A	12678		897	MSNRVVCREASHAGSWYTASG PQLNAQLEGWLSQVQSTKRPA RAIIAPHAGYTYCGSCAAHAYK QVDPSITRRIFILGPSHHVPLSRC ALSSVDIYRTPLYDLRIDQKIYG ELWKTGMFERMSLQTDEDEHS IEMHLPYTAKAMESHKDEFTIIP VLVGALSESKEQEFGKLFSKYL ADPSNLFVVSSDFCHWGQRFR Y\SYYDESQGEIY*SIEHLDKMG MSIIEQLDPVSFSNYLKKYHNTI CGRHPIG\VLLNAITELQKNGM NMSFS\FLNYAQSSQC\RNWQD SSVSYAAGALTVH
12603	42971	A	12679	1	912	
12604	42972	A	12680 12681	112	156 1120	
12605	42973 42974	A	12682	33	415	NMNKPITPSTYVRCLNVGLIRK LSDFIDPQEGWKKLAVAIKKPS GDDRYNQFHI\DAVPKTANTLP SKEAITVQQKQMPFCDKDRTL MTPVQNLEQSYMPPDSSSPENK SLEVSDTRFHSFSFYELKN
12607	42975	A	12683	3	311	
12608	42976	Α	12684	1	1125	
12609	42977	Α	12685	1	1365	
12610	42978	A	12686	3	141	VVSTVVPDS/AHKLFIGGLPNYL NDDQCHGLSKGYAFCEYVDIN VTV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	1
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12611	42979	A	12687	1	1255	GTRSTSSLRRQMKNIVHNYSEA
				1		EIKVREATSNDPWGPSSSLMSEI
						ADLTYNVVAFSEIMSMIWKRL
						NDHGKNWRHVYKAMTLMEYL
						IKTGSERVSQQCKENMYAVQT
						LKDFQYVDRDGKDQGVNVRE
						KAKQLVALLRDEDRLREERAH
						ALKTKEKLAQTATASSAAVGS
ŀ						GPPPEAEQAWPQSSGEEELQLQ
						LALAMSKEEADQEERIRRGDDL
			:	1		R\RRWQSRRARGRLGARRSRPS
		ł				WTLLTSSRPQLLPRPQTPGGAQ
						HPWLLPSPRLPPPRTPGAAPLSL
		1				
		ł		-		QLLIPGEVQPPRRPLGTPGGLLP
				}		LQDPQLTLGVGPQPLQLGRGPR
						LIHGEVPMVGSRSVGPQPPIPGH
				1		RPRPSQIPGEGHLPSPAPMAQQ
						QPGDSTRSPTSSLTLTDSARHCR
	_					PPGAAQESWSCWQERCRPEAL
12612	42980	A	12688	116	1541	
12613	42981	A	12689	3	415	
12614	42982	Α	12690	1	1440	
12615	42983	C	12691	117	314	
12616	42984	A	12692	3	120	
12617	42985	A	12693	1	729	
12618	42986	Α	12694	79	870	EWSSVRRSLVEKRALRRPHPQC
						LCFRMKTILSNQTCRPFPE\NVD
			[\ITLKGRP\VIV*GPAEGTLR\RDF
						\NHLQW*NLSLSWKEKKRGSR
						VDK\WWGNRKGNWPT\VRTYL
						VVHVQN\MIKGVLPLGLPVTKD
						EGLVYA\HFPHPTLVIPGRNGSS
	F	ļ				LLKSRNFLGVKNTSRRVSG*RP
	:					G\VACS\VSSSPRKDEFNPLKGN
					İ	DIGACFQISAGFGFQQAPHTCL
						KTRDIQGNFWDGYPMVLWKK
						GTVSARAGWNKDLKEFTWATE
	į				,	RKMPG
12619	42987	A	12695	2	293	
12620	42988	Α	12696	1	273	TRGPWCDSVLRGCSLEQRSFIS
						VRLLSYLSACRHPMEDSMDMD
						MSPLRPQNYLFG\SLGAGAKDE
				1		LHIVEAEAMNYEGSPIKVTLAT
			l			LKMS
12621	42989	Α	12697	1	1308	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon,/=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible πucleotide insertion)
12622	42990	Α	12698	124	1135	DKNSMGLPGVIPSCAAVLWSSI
						HLSSSALSPTISACRHPMEDSM
						DMDMSPLRPQNYLFGCELKAD
						KDYHFKVDNDENEHQLSLRTV
						SLGAGAKDELHIVEAEAMNYE
						GSPIKVTLATLKMSVQPTVSLG
	i		ļ			GFEITPPVVLRLKCGSGPVHISG
						QHLVAVEEDAESEDEEEEDVK\
						LLSISGKRSAPGGGSKVPQKK\V
						KLAAD\EDDDD\DDEEDDDEDD
						\DDDDF\DDEEAEEKAPVK\KSI\
						RDTPAKNAQKSNQNGKDSKPS
						STPRSKGQESF\KKQEKTPKTPK
						GPSSVEDIKAKMQASIA/EKGGS
						LPKVEAKFINYVKNCFRMTDQ
12622	42001	ļ_	12600	201	416	EAIQDLWQWRKSL
12623 12624	42991 42992	A	12699 12700	201	416 678	
12625	42993	A	12701	3	2149	
12626	42994	A	12701	1	219	
12627	42995	A	12704	267	437	
12628	42996	A	12705	507	845	
12629	42997	A	12706	426	704	SRRWGGVHSCPPSPPPTAGHWT
			1			LLILCAKARRSEAAMVLASRRR
						RPGPRPRRRRRLPVLGSRGGAG
						PARGRGPSGWPGPLRAAPSP*P
						SSTV
12630	42998	A	12707	195	892	
12631	42999	A	12708	1	893	
12632	43000	A	12709	1	777	WEOV A DOTTO CLODE DOCUMENTOR
12633	43001	A	12710	1	439	KTQVAPSTDGIQP\PSSNSRTDE
						REFFIASYNRKKEDGEGNVWIA
		1				KSSAGAKGEGILISSEASELLDFI DNQAQVHVIQKYLEHPLLLEPG
						HRKFDIRSWVLVDHQYNIYLY
						REGVLRTASEPYHVDNFQDKT
						CHLTNHCIQKKKKK
12634	43002	A	12711	1	394	SGTRPVFLVPHTIGIP\HPAIVTP
12054	13002	1.	12/11	1		QVKQEHPHTDSDLMHVKPQHE
						QRKEQEPKRPHIKKPLNAFMLY
						MKEMRANVVAECTLKESAAIN
						QILGRRWHALSREEQAKYYEL
						ARKERQLHMQLYPGWSARDN
12635	43003	A	12712	1	789	
12636	43004	A	12713	1	870	
12637	43005	Α	12714	1	729	
12638	43006	Α	12715	1	828	
12639	43007	Α	12716	1	213	
12640	43008	Α	12717	1	664	
12641	43009	A	12718	1	695	
12642	43010	A	12719	96	269	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12643	43011	A	12720	3	241	
12644	43012	A	12721	213	660	EGPARHRLLSPVRASKMTKKR
12011	13012	•	12,2,			RNNGRAQKRAARHVRGPSPLK/
						CFPSGPLPPNCAR\CVPK\DKAIK
						KF\VI\RNIVEAAA\AR\DISEASV
						FDAYL\LPKL\YVKLH\YCVSCAI
						HKQK*SGNRSS*SPAKTRTPPPR
						FR\PGG*LPHGPPTKSP
12645	43013	A	12722	1	303	
12646	43014	Α	12723	610	928	FLSLPTFLFVIFSGEEELLVLALV
						FLSLFFFFFFLRWSFAVVAQAV\
						VQWHNLSSLPFGFKQFSCLSLP
						SSWDYRCPPPRPANFCIFTRDG
						VSPCCPGWSRTSDLR
12647	43015	A	12724	1	2775	
12648	43016	Α	12725	1	6039	
12649	43017	Α	12726	2	3661	
12650	43018	A	12727	1	1173	
12651	43019	Α	12728	1	1632	MPNPKNSKGGRKNKRANSSGD
						EQENGAGALAAAGAAGAAAG
1						GALAAAAGCGAAAAGAPGAG
}						GAAGAGGAGTGAANAAAAG
1						AAAAGDAKNEAPCATPLICSFG
						RPVDLE\KDDYQKVVCNNEHC
						PCSTWMHLQCFYEWESSILVQF
						NCIGRARTWNEKQCRQNMWT
						KKG\YDL\AFRFCSCRCGQGNL
						KKDTDWYQVKRMQDEKKKKS
						GSEKNTGRPPGEAAEEAKKCRP
						PNKPQKGPSHDLPRRHSMDRQ
						NSQEKAVGAAAYGARSPGGSP
						GQSPPTGYSILSPAHFSGPRSSR
						YLGEFLKNAIHLEPHKKAMAG
						GHVFRNAHFDYSPAGLAVHRG
1						GHFDTPVQFLRRLDLSELLTHIP RHKLNTFHVRMEDDAQVGQG
1						,
						EDLRKFILAALSASHRNVVNCA
						LCHRALPVFEQFPLVDGTLFLSP SRHDEIEYDVPCHLQGRLMHL
						YAVCVDCLEGVHKIICIKCKSR
						WDGSWHQLGTMYTYDILAASP
İ						CCQARLNCKHCGKPVIDVRIG
						MQYFSEYSNVQQCPHCGNLDY
						HFVKPFSSFKVLEAY
12652	43020	A	12729	1	1485	THE COLUMN DESTRICT
12653	43021	A	12730	1	1074	
12654	43022	Α	12731	2	3394	
12655	43023	Α	12732	1	2190	
12656	43024	A	12733	1	1251	
12657	43025	Α	12734	1	1710	
12658	43026	Α	12735	3	679	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12659	43027	A	12736	209	337	
12660	43028	Α	12737	365	538	
12661	43029	Α	12738	2	196	
12662	43030	Α	12739	I	2639	
12663	43031	Α	12740	3	173	
12664	43032	Α	12741	1	1479	
12665	43033	Α	12742	1	921	
12666	43034	Α	12743	1	798	
12667	43035	A	12744	2	773	
12668	43036	В	12745	75	321	
12669	43037	A	12746	321	738	
12670	43038	Α	12747	1	3190	
12671	43039	Α	12748	287	1636	ACGAMERAPPDGPLNASGALA
						GDAAAAGGARGFSAAWTAVL
	•					AALMALLIVATVLGNALVMLA
			i			FVADSSLRTQNNFFLLNLAISDF
						LVGAFCIPLYVPYVLTGRWTFG
	1					RGLCKLWLVVDYLLCTSSAFNI
						VLISYDRFLSVTRAVSYRAQQG
						DTRRAVRKMLLVWVLAFLLYG
		1				PAILSWEYLSGGSSIPEGHCYAE
					:	FFYNWYFLITASTLEFFTPFLSV
						TFFNLSIYLNIQRRTRLRLDGAR
						EAAGPEPPPEAQPSPPPPPGCWG
						CWQKGHGEAMPLHRYGVGEA
						AVGAEAGEATLGGGGGGGSVA
						SPTSSSGSSSRGTERPRSLKRGS
						KPSASSASLEKRMKMVSQSFTQ
						RFRLA\RDRKEAKSLAVIVSIFG
						LCWAPYTLLMIIRAACHGHCVP
						DYWYETSFWLLWANSAVNPV
						LYPLCHHSFRRAFTKLLCPQKL
10.650	10010	 	10740		0.140	KIQPHSSLEHCWK
12672	43040	A	12749	11	2142	THURSDAY MANDETTEL DETAIL
12673	43041	A	12750	3	711	VWWNSDLMVINRSTTELPLTV
						SYDKVSLGRLRFWIHMQDAVY SLQQFGFSEKDADEVKGIFVDT
						NLYFLALTFFVAAFHLLFDFLA
						FKNDISFWKKKKSMIGMSTKA VLWRCFSTVVIFLFLLDEQTSLL
			ļ			VLVPAGVW/AAAIELWES*RK/
	1					VI*RWTIFWRGLMPEFQFGTYS
						-
						ESERKTEEYDTQAMKYLSYLL YPLCVGGAVYSLLNIKYKSWY
						SWLINSFVNGVYAFGFLFMLPQ
12674	43042	A	12751	1	2025	PARTIMOLANDA LALOLITAIDA
12674	43042	A	12752	494	747	ATVKWITLCRNRNIKLPTFIKKE
120/3	CFUCE	^	12/32	77	171	GTGKAPKKSRKSVLGLGP*VHT
						SPGDPAEGEGLPAGERPDGNLS
				1		
12676	43044		12753	9	731	ZOLINGDIG LIKKKINITA
12676	43044	A	12753	9	731	QSPNSLRFYIKKKKAYK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12677	43045	A	12754	143	641	RSWVLLGVGARGSTPRTYQVP
						SPSLAPSALC*VS*VPLESLSCSH
						ADNWKQELTKFISPDQLPVEFG
						GTMTDPDGNPKCLTKINYGGE
						VPKSYYLCKQVRLQYEHTRSV
						GRGSSLQVENEILFPGCGLR\VL
						RFYNTYSLVHSKRISYTVEVLL
						PDQTFMEKMEKF
12678	43046	Α	12755	1	1506	
12679	43047	Α	12756	2	526	CQPVRAA*RHGNTPLATV*PAS
						T/WWKWSEKQKEEQEMQNWR
						ESAS/VSGYPQDILSCAMSQNPR
					:	KRKRPRNVSAPPVHQMAQMNI
						SVEQMETSKTHQATPLTWGQM
						KRLAHVAEENLRSQNKPLTTSN
						LMVAMMVVISLAVSLPVAGAD
						QNYIYWAYIPFPH
12680	43048	Α	12757	1	1443	
12681	43049	Λ	12758	3	766	
12682	43050	A	12759	29	253	
12683	43051	A	12760	34	252	COM HIDING A DI MONELLA COMMUNI
12684	43052	A	12761	1362	1611	SSVLHPWDARLVQHT\KSIHIIH
						HINRTNDKNHMIISIDAEKAFD
						KIQQDFMLKTLNKLGIDGTYLK
12685	43053	В	12762	1	1755	IVRATYDKPTASIILNG
12686	43054	A	12763	1	2109	
12687	43055	В	12764	1	1497	
12688	43056	В	12765	1	1527	
12689	43057	В	12766	1	1746	
12690	43058	В	12767	1	2853	
12691	43059	В	12768	1	1479	
12692	43060	В	12769	1	1932	
12693	43061	A	12770	1269	1581	NTGKPNPAAHQKGYPPRSSGPH
						PWDARLVQHT\KSINIIHHINRT
						KDKNHMIISIDAEKAFDKIQLPF
						MLKTLSKLGVSYNMFTRLTIPI
						AQVDTRGNQLPSYLI
12694	43062	В	12771	1	4404	
12695	43063	В	12772	56	2080	
12696	43064	В	12773	1	5445	
12697	43065	В	12774	295	2831	
12698	43066	Α	12775	89	459	
12699	43067	A	12776	2	980	
12700	43068	A	12777	1	378	
12701	43069	Α	12778	1	363	PGVALPGRCRQAPADLLP*MR
						AYWPDVIYSFANRSRFWKHEW
				1		EKHGTCAAQVDALNSQNKYFG
		1				TSLELYRELDLNSVLLKLGIKPS
		1				INYYQVADFKDALARVYGVIP
L	<u> </u>	<u> </u>		<u>L </u>		KIQCLPPSQDEDRQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12702	43070	A	12779	1	577	QRFKNDCRDPRDTGPIHGLWPD
						KSEG/CNRSW/PFNLEEIKDLCP
						EMRAYWPDVIHSFPNRSRFWK
						HEWEKHGTCAAQVDALNSQK
						KYFGRSLELYRELDL/NQVADF
	1					KDALARVYGVIPKIQCLPPSQD
	ļ					EEVQTIGQIELCLTKQDQQLQN
						CTEPGEQPSPKQEVWLANGAA
						ESRGLRVCEDGPVFYPPPKKTK
12703	43071	A	12780	1	624	MRPAALRGALLGCLCLALLCL
12703	15071	` `	1.2,00		021	GGADKRLRASVSRPFSLEHPEP
						GHCLDLHLRAYPAR\WSS*PWP
						QDFWVR*PTWES/VSDLLPEMR
						AYWPDVIHSFPNRSRFWKHEW
	ŀ					EKHGTCAAQVDALNSQKKYFG
						RSLELYRELDLNRGERVPTQFP
	1					RELVVNRSLAPPPTPGFLSGHVI
						SVHTSSPLPSAMSGSFLRPLPEA
						ESGSMLLGQPADP
12704	43072	A	12781	1	612	EBGSMEEGQI NEI
12705	43073	A	12782	113	1149	RGAPRGPGRQGCCGHSRSPAR
12703	1.5075	'	12702			GPRDTGLGRPRAPGAPGGRCCS
						GGSAAARGLGDSVSPGSLPRAA
						GGKWEPERQLAAFSPQVGTMR
						PAALRGALLGCLCLALLCLGG
						ADKRLRDNHEWKKLIMVQHW
						PETVCEKIQNDCRDPPDYWTIH
						GLWPDKSEGCNRSWPF\NLEEI
						KDLLPEMRAYWPDVIHSFPNRS
				= -		RFWKHE/WPKRHGTCAAQ\VD
		1	}			ALNSQK\KYFGRSLELY\RKLDL
		ŀ				NSVL\LKLGIKPSINYYQVADFK
						DALARVYGVIPKIQCLPTSQGE\
						EVQTIGQ\IDLCLIKARTRQLQN\
						CTEPGEQP\SPKQEVWLANGA\
						AESRGLRVCEDGPVFYPPPKKT
12706	43074	A	12783	2	515	TESTOSKY CESSI VI IIII KKI
12707	43075	В	12784	62	359	
12708	43076	В	12785	219	523	
12709	43077	A	12786	1010	1550	DTENIPSSRRRERSKVPYIVRQC
		1				VEEIERRGMEEVGIYHVSGR\V
	}					AADIQALKAAFNVSECRPAQD
						GMEVWAVVSAMRSQSAPRPRH
						VTSFSVSFLHLSGSSRRPLHFRA
	1					LS\NNKDVSVMMSEMDVNAIA
		1				GSLKLYFRELPEPLFTDEFYPNF
[AEGIGEHWRPWPHGRRLLHVH
						CCPQRL
	<u> </u>		<u> </u>	L		1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12710	43078	Α	12787	2	951	GSNLYCTLEGDSFGYFVNKAK
						TRVYRD/THADPNWNEEFEIEL
		İ				EGSQTLRILCYEKCYNKTKIPKE
1						DGESTDRLMGKGQVQLDPQAL
						QDRDWQRTVIAMNGIEVKLSV
		l				KFNSREFSLKRMPSRKQTGVFG
]				VKIAVVTKRERSKV\PYI\VRQC
		i				VEEIEPPKALEGGWASTACPAG
ŀ						VATDIQALKAAFDGNNKDVSV
						MMSEMDVNAIAGTLNLYFREL
	İ					PEPLFT\DEFYPNFAEG/IRVAEK
						EAVNNM\SLHNLATVFGPTLLR
						PSEKESKLPANPS\QPITMTDSW
						SLEVMSQVQVLLYFLQLEAIPA
:						PDSKRQSILFSTEV
12711	43079	A	12788	3	644	
12712	43080	Α	12789	168	378	
12713	43081	Α	12790	1	486	
12714	43082	Α	12792	83	536	
12715	43083	Α	12793	1	662	
12716	43084	Α	12794	3	357	
12717	43085	A	12795	2	421	
12718	43086	Α	12796	2	908	
12719	43087	Α	12797	1	846	
12720	43088	C	12798	87	413	
12721	43089	Α	12799	1	508	MCQGDTYAWGGHMCRGGHLR
						RGHSHERDTPRGPERPPRPPAR
						DRESKDERRRPPPKDPPAVRTC\
						PDSPSTATPPPPPPPPPPPPPPPPP
						PAAPSAFAALRLRGPGRKWTA
						RGQLPLPATAPEARVRALPRGG
						YSSPGRATSARGPAPWPLPSRT
						GKPPRPCSANTQEHC
12722	43090	Α	12800	1	393	
12723	43091	Α	12801	1	582	
12724	43092	Α	12802	1	1422	
12725	43093	A	12803	1	1080	0.0000.00100000000000000000000000000000
12726	43094	A	12804	2	261	SCPFGGNCFYKHAYPDGRREEP
						QRQKVGTSSRYRAQRRNHFWE
		İ				LIEERENSNPFDNDEEEVVTFEL
10707	12005	_	12007	60	1000	GE\MLLMLLAAGGDDELTNS
12727	43095	В	12805	60	802	<u> </u>
12728	43096	A	12806	2	238	
12729	43097	A	12807	1	1017	
12730	43098	В	12808	1	939	
12731	43099	Α	12809	11	708	

SEQ ID		Met	SEQ ID NO:	Nucleotide	1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
12732	43100	A	12810	1	2653	MGDFNTPLSTLDRSMRQKVNK
						DIQELNSALHQADLIDIYRNLHP
						ESTEYTFFSAPHHTYSKIDHILG
						SKAPLSKYRRSEIKINCLSDHSA
						IKLELRIKKLTQNRSTTWKLNN
						LLLNDYWVHNEMKAEIKMFFE
	!		į			TNENKDTTYQNLWDTLKAVCR
	!					GKFIALNAHKRKQERSKIDTLT
						SQLKELEKQEQTHSKASRRQEI
	l i					SKIRGELKEIETQKTLQKINESR
						GWFFEKINKIDRLLARLIKKKR
				:		EKNQIHAIKNDKGDMSTNHTEI
						QTTIREYYKHLYANKLENLKEI
						DKFLETYSLPRLNQEEVESLNR
				;		PITGSEIEAIINSLPNKRSPGPDG
						FTAKFYQRYKEELLISNFSKVS
	İ					GYKINVQKSQAFLYTNNRQTES
1					}	QIMSELPFTIASKRMKYLGIQPT
						RDMKDLFKENYKPLLNEIKEDT
						NKWKNIPCSWVGRINIVKMAIL
						PKNWKKTTLKFIWNQKRARIA
						KSILSQKNKAGGIMLPDFKLYY
ĺ	•					KATITKTAWYWYQNRDIDQW
1						NRTEPSEIMPHIYNHLIFDKPDK
ĺ						NKKWGKDSLFNKWCWENWLA
	<u> </u>					ICRKLKLDPFLTPYTKINSRWIK
						DLNVRPKTIKTLEENPGNTIQDI
						GMGEDFMSKTPEAMATKAKID
						KWDLIKLKSFCTAKETTIRLNR
				,		PPTEWEKIFAIYSSDKGLISRIYN
						ELQQIYEKKTNNPIKKWAKDM
		l				NRHLSKEDIYAAKRHMKKCAS
12733	43101	В	12811	1	2748	
12734	43102	Α	12812	1	1962	
12735	43103	В	12813	1	1833	
12736	43104	Α	12814	176	421	QQPASPTRSVLLFPKTAHISGSP
						APPNVGLLNPSTGAAQKKTMT
						SAGLGRLSL*K*LNLPECVACA
<u></u>						GETWVTSMMSGNQCN

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12737	43105	A	12815	1	1615	LISNFSKVSGYKINVQKSQAFL
ļ						YTNNRQTESQIMSELPFTIASKR
1						MKYLGIQPTRDMKDLFKENYK
						PLLNEIKEDTNKWKNIPCSWVG
					1	RINIVKMAILPKNWKKTTLKFI
						WNQKRARIAKSILSQKNKAGGI
						MLPDFKLYYKATITKTAWYWY
						QNRDIDQWNRTEPSEIMPHIYN
						HLIFDKPDKNKKWGKDSLFNK
						WCWENWLAICRKLKLDPFLTP
						YTKINSRWIKDLNVRPKTIKTLE
ŀ						ENPGNTIQDIGMGEDFMSKTPE
ļ						AMATKAKIDKWDLIKLKSFCT
						AKETTIRLNRPPTEWEKIFAIYS
						SDKGLISRIYNELQQIYEKKTNN
						PIKKWAKDMNRHLSKEDIYAA
						KRHMKKCASSLAIREIQIKTTM
						RYHLTPVRMAIIKKSGNNRCW
j						RGCGEIGTLLHCWWDCKLVQP
						LWKSVWRFLRDLELEIPFDPAIP
						LLGIYPKDYKSCCYEDTCTHMF
						IVALFTIAKTWNQPKCPTMIDW
						IKKMWHIYTMEYYADIKKDEF
						MSFVRTRMKLETIILSKISQ\EK
						KTKHRMFSLIGGN
12738	43106	A	12816	1	996	
12739	43107	Α	12817	1	756	
12740	43108	Α	12818	1	1428	

1629

12741 43109 A 12819 1 2377	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
12741 43109 A 12819 I 2377	NO:	1	hod		ľ	2	*=Stop codon, /=possible nucleotide
12741 43109 A 12819 1 2377		sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
IHHDQVGFIPGMQGWFNIRKSI NIVQHINRTNDKNHMISIDGIDG AFDRIQQFMLKTLNKLGIDG YLKIIRAIYDKSTANIVLNGQK EAFPLKTGTRQGCPLSPLLTIMI LEVLARAIRQF&EIKGIQLRKE IKLSLFADDMIVHLENPIVSAQ LLKLIDNFSKVSGYKINVQKS AFLYTNNRQTESQIMSELPFTI SERIKYLGIQLTRDVKDLFKEN YKPLLNDIREDTNKWKNTPRS WVGRINIMKMAILPKVIYRFN, IPIKLPMTFFTELEKTTLKFIWN QKGARIAKSILIYRFNAQELEK TLKFIWNQKGARIAKSILIYRFNAQELEK TLKFIWNQKGARIAKSILSQKN KAGGITLPDFKLYYKATVIKTA WYWQNRIDQWNIXTEPSIL PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKLKLD FLIPYTKINSRWIKDLNVRPKTI KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDK WDLINLK SFCTAKETTIRVNRQPIEWEBI AIYSSDKGRISTYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGIN DWCWRGGGIGRLLHCWWD KLVQPLWKSVWRFLRDLELET PFDPATPLLGIYPKDYKSCYCL DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES* RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH					sequence		
NIVQHINRTNDKNHMIISIDAEI AFDKIQQRFMLKTLNKKIGIDG YLKIIRAIYDKSTANIVLNOGK EAFPLKTGTRQCPLSPLLFNIV LEVLARAIRQEKEIKGIQLRKE IKLSLFADDMIVHLENPIVSAQ LLKLIDNFSKVSGYKINVQKSC AFLYTNNRQTESQIMSELPFTIV SERIKYLGIQLTROVKOLFKEN YKPLLNDIKEDTNKWKNTPRS WVGRINIMKMAILPKVIYRFN, IPIKLPMTFFTELEKTTLKFIWN QKGARIAKSILIYRFNAQELEK TLKFIWNQKGARIAKSILSQKN KAGGITLPDFKLYYKATVIKTA WYWYONRDIDQWNRTEPSEIB PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKLKLD FLIPYTKINSRWIKDLNVRPKTI KTLEENLGSTTQDIAMGKDFM KTPKAMATKAKIDKWDLINLK SFCTAKETTIRVNRQPIEWEKI AAIYSSDKGRISRIYNELGVIYKK KTNNPIK WAKDMNRHFSKEI IYAAKRHMKKCSSLAIREMH KTTIMRIHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDC KLVQPL WKSVWRFLENDLELET PPDPATPLLGIYPKDYKSCCTY DICTRVCVPAALFTIANTWNP 12742 43110 A 12820 I 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES* RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH	12741	43109	Α	12819	1	2377	MNIDAKILNKILANRIQQHIKKL
AFDKIQQRFMLKTLNKLGIDG			ł				IHHDQVGFIPGMQGWFNIRKSI
AFDKIQQRFMLKTLNKLGIDG							NIVQHINRTNDKNHMIISIDAEK
EAFPLKTGTRQGCPLSPLLFNIN LEVLARAIRQEKEIKGIQLRKE IKLSLFADDMIVHLENPIVSAQ LLKLIDNFSVSGYKINVQKSQ AFLYTNNRQTESQIMSELPFTIL SERIKYLGIQLTRUVDLFKEN YKPLLNDIKEDTNKWKNTPRS WVGRINIMKMAILPKVIYRFN/ IPIKLPMTFFTELEKTTLKFIWN QKGARIAKSILISQKN KAGGITLPDFKLYYKATVIKTA WYWYQNRDIDQWNTEPSEIN PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKLKLD FLIPYTKINSRWIKDLNVRFKTI KTLEENLGNTIQDIAMGKDFM KTPKAMATAKIDKWDLINLR SFCTAKETTIRVNRQPIEWEKII AIYSSDKGRISRIYMELQTIYK KTNPIKKWAKDMNRHFSKEI IYAAKRIMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLHCWWDG KLVQPLWKSVWRFLRDLELET PFDPATP\LLGIYPKDYKSCCYP DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*C RGIQKKASLRSRTONHSASSGI ESRV*GLGTTGYRGTASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH	ļ						AFDKIQQRFMLKTLNKLGIDGT
LEVLARAIRQEKEIKGIQLRKE IILSLFADDMIVHLENPIVSAQ LLKLIDNFSKVSGYKINVQKSC AFLYTNIRQTESQIMSELPFTIL SERIKYLGIQLTRDVKDLFKEN YKPLLNDIKEDTINKWNTPRS WVGRINIMKMAILPKVIYRFN, IPIKLPMTEFTELEKTTLKFIWN QKGARIAKSILIYRFNAQELEK TLKFIWNQKGARIAKSILSQKN KAGGITLPDFKLYYKATVIKTA WYWQNRDIDQWNRTEPSEIN PHIYNHLIFDKPDKNKKWGND LFNKWCWENVLALGIKLKLD FLIPYTKINSRWIKDLNVRPKTI KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINLR SFCTAKETTIRVNRQPIEWEKII AIYSSDKGISRIYNELKQIYKR KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKLGN DWCWRGGEIGRLLHCWWD KLVQPLWKSVWRFLRDLELET PFDPATPILLGIYPKDYKSCCYI DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES* RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							YLKIIRAIYDKSTANIVLNGQKL
IKLSLFADDMIVHLENPIVSAQ							EAFPLKTGTRQGCPLSPLLFNIV
IKLSLFADDMIVHLENPIVSAQ							LEVLARAIRQEKEIKGIQLRKEE
AFLYTNNRQTESQIMSELPFTIA SERIKYLGIQLTRDVKDLFKER YKPLLNDIKEDTNKWKNTPRS WYGRIMKMAILPKVIYRFN. IPIKLPMTFFTELEKTTLKFIWN QKGARIAKSILIYRFNAQELEK TLKFIWNQKGARIAKSILSQKN KAGGITLPDFKLYYKATVIKTA WYWYQNRDIDQWNRTEPSEIN PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKUKLD FLIPYTKINSRWIKIDLNVRPKT KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINLK SFCTAKETTIRVNRQPIEWEKII AIYSSDKGRISRIYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLHCWWDD KUVQPLWKSVWFLRDLELET PFDPATPLLGIYPKDYKSCCYP DICT/RVCVPAALFTIANTWNP TOTAL TO							IKLSLFADDMIVHLENPIVSAQN
AFLYTNNRQTESQIMSELPFTIA SERIKYLGIQLTRDVKDLFKER YKPLLNDIKEDTNKWKNTPRS WYGRIMKMAILPKVIYRFN. IPIKLPMTFFTELEKTTLKFIWN QKGARIAKSILIYRFNAQELEK TLKFIWNQKGARIAKSILSQKN KAGGITLPDFKLYYKATVIKTA WYWYQNRDIDQWNRTEPSEIN PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKUKLD FLIPYTKINSRWIKIDLNVRPKT KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINLK SFCTAKETTIRVNRQPIEWEKII AIYSSDKGRISRIYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLHCWWDD KUVQPLWKSVWFLRDLELET PFDPATPLLGIYPKDYKSCCYP DICT/RVCVPAALFTIANTWNP TOTAL TO							LLKLIDNFSKVSGYKINVQKSQ
SERIKYLGIQLTRDVKDLFKEN YKPLNDIKEDTNKWKNTPRS WVGRINIMKMAILPKVIYRFN, IPIKLPMTFTFELEKTTLKFIWN QKGARIAKSILIYRFNAQELEK TLKFIWNQKGARIAKSILSQKN KAGGITLPDFKLYYKATVIKT WYWYQNRDIDQWNRTEPSEIR PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKLKLD FLIPYTKINSRWIKDLNVRRKTI KTILEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINLK SFCTAKETTIRVNRQPIEWEKIF AIYSSDKGRISRIVNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDG KLVQPLWKSVWRFLRDLELET PFDPATPILLGIYPKDYKSCCYP DICT/RVCVPAALFTIANTWNP 12742							AFLYTNNRQTESQIMSELPFTIA
WVGRINIMKMAILPKVIYRFN/ IPIKLPMTFFTELEKTTLKFIWN QKGARIAKSILYRFNAQELEK TLKFIWNQKGARIAKSILSQKN KAGGTLPDFKLYYKATVIKTY WYWYQNRDIDQWNRTEPSEIN PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKLKLD FLIPYTKINSRWIKDLNVRPKTI KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINLK SFCTAKETTIRVNQPIEWEKI AIYSSDKGRISRIYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGGGEIGRLLHCWWDC KLVQPLWKSVWRFLRDLELET PFDPATPILIGIYPKDYKSCCYY DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*C RGIQKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							SERIKYLGIQLTRDVKDLFKEN
IPIKLPMTFFTELEKTTLKFIWN QKGARIAKSILIYRFNAQELEK TLKFIWNQKGARIAKSILSQKM KAGGITLPDFKLYYKATVIKTA WYWYQNRDIDQWDRTEPSEIN PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKLKLD FLIPYTKINSRWIKDLNVRPKTI KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDK WDLINLK SFCTAKETTIRVNRQPIEWEKI AIYSSDKGRISRIYNELKQIYKK KTNNPIKK WAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDC KLVQPLWKSVWRFLRDLELET PFDPATP\LLGIYPKDYKSCCYFDICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840							YKPLLNDIKEDTNKWKNTPRS
QKGARIAKSILIYRFNAQELEK							WVGRINIMKMAILPKVIYRFNA
TIKFIWNQKGARIAKSILSQKN KAGGITLPDFKLYYKATVIKTA WYWQNRDIDQWNRTEPSEII PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKLKLD FLIPYTKINSRWIKDLNVRPKTI KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINLK SFCTAKETTIRVNRQPIEWEKIF AIYSSOKGRISRIYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDC KLVQPLWKSVWRFLRDLELET PFDPATP\LLGIYPKDYKSCCYF DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840]						IPIKLPMTFFTELEKTTLKFIWN
KAGGITLPDFKLYYKATVIKTA WYWYQNRDIDQWNRTEPSEIN PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKLKLD FLIPYTKINSRWIKDLNVRPKTI KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINI.K SFCTAKETTIRVNRQPIEWEKII AIYSSDKGRISRIYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDC KLVQPLWKSVWRFLRDLELET PFDPATPLLGIYPKDYKSCCYI DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*C RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH					ļ		QKGARIAKSILIYRFNAQELEKT
KAGGITLPDFKLYYKATVIKTA WYWYQNRDIDQWNRTEPSEIN PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKLKLD FLIPYTKINSRWIKDLNVRPKTI KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINI.K SFCTAKETTIRVNRQPIEWEKII AIYSSDKGRISRIYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDC KLVQPLWKSVWRFLRDLELET PFDPATPLLGIYPKDYKSCCYI DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*C RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							TLKFIWNQKGARIAKSILSQKN
PHIYNHLIFDKPDKNKKWGND LFNKWCWENWLAICRKLKLD FLIPYTKINSRWIKDLNVRPKTI KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINLK SFCTAKETTIRVNRQPIEWEKIH AIYSSDKGRISRIYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDG KLVQPLWKSVWRFLRDLELET PFDPATPULLGIYPKDYKSCCYF DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*G RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							KAGGITLPDFKLYYKATVIKTA
LFNKWCWENWLAICRKLKLD FLIPYTKINSRWIKDLNVRPKTI KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINLK SFCTAKETTIRVNRQPIEWEKIF AIYSSDKGRISRIYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDG KLVQPLWKSVWRFLRDLELET PFDPATPVLLGIYPKDYKSCCYF DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*C RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH			İ				WYWYQNRDIDQWNRTEPSEIM
FLIPYTKINSRWIKDLNVRPKTI KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINLK SFCTAKETTIRVNRQPIEWEKIF AIYSSDKGRISRIYNELKQIYKK KTNNPIKK WAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGGEIGRLLHCWWDC KLVQPLWKSVWRFLRDLELET PFDPATP\LLGIYPKDYKSCCYF DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 EYKNR*GLSLNPWGKTVHISW DVWSVGSKAKRNWESWES*CRIOCK GIOKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							PHIYNHLIFDKPDKNKKWGNDS
KTLEENLGNTIQDIAMGKDFM KTPKAMATKAKIDKWDLINLK SFCTAKETTIRVNRQPIEWEKIF AIYSSDKGRISRIYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDG KLVQPLWKSVWRFLRDLELET PFDPATP\LLGIYPKDYKSCCYF DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*G RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPT HMSQLMWTACPKGLGIALICL VRH					1		LFNKWCWENWLAICRKLKLDP
KTPKAMATKAKIDK WDLINLK SFCTAKETTIRVNRQPIEWEKIF AIYSSDKGRISRIYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDC KLVQPLWKSV WRFLRDLELET PFDPATP\LLGIYPKDYKSCCYF DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 I 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*GRIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTH HMSQLMWTACPKGLGIALICL VRH							FLIPYTKINSRWIKDLNVRPKTI
SFCTAKETTIRVNRQPIEWEKIF							KTLEENLGNTIQDIAMGKDFMS
AIYSSDKGRISRIYNELKQIYKK KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDG KLVQPLWKSVWRFLRDLELET PFDPATP\LLGIYPKDYKSCCYRDICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*GRIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							KTPKAMATKAKIDKWDLINLK
KTNNPIKKWAKDMNRHFSKEI IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDC KLVQPLWKSVWRFLRDLELET PFDPATP\LLGIYPKDYKSCCYP DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 l 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*(RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							SFCTAKETTIRVNRQPIEWEKIF
IYAAKRHMKKCSSSLAIREMH KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDC KLVQPLWKSVWRFLRDLELET PFDPATP\LLGIYPKDYKSCCYR DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 l 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*GRGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							AIYSSDKGRISRIYNELKQIYKK
KTTMRHHLTSVRMAIIKKLGN DWCWRGCGEIGRLLHCWWDC KLVQPLWKSVWRFLRDLELET PFDPATP\LLGIYPKDYKSCCYP DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*C RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH			İ				KTNNPIKKWAKDMNRHFSKED
DWCWRGCGEIGRLLHCWWDCKLVQPLWKSVWRFLRDLELETPFDPATP\LLGIYPKDYKSCCYRDICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 I 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISWDVWSVGSSKAKRNWESWES*GRGIQKKASLRSRTGNHSASSGIESRV*GLGTTGYRGITASLMSLRSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSDQTLVVQRGQKMEQANHPDPTHHMSQLMWTACPKGLGIALICLVRH			İ				IYAAKRHMKKCSSSLAIREMHI
KLVQPLWKSVWRFLRDLELET PFDPATP\LLGIYPKDYKSCCYR DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 1 4840 12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*GRIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							KTTMRHHLTSVRMAIIKKLGN
PFDPATP\LLGIYPKDYKSCCYRDICT/RVCVPAALFTIANTWNP							DWCWRGCGEIGRLLHCWWDC
DICT/RVCVPAALFTIANTWNP 12742 43110 A 12820 I 4840 484111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*6 RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPT HMSQLMWTACPKGLGIALICL VRH							KLVQPLWKSVWRFLRDLELET
12742 43110 A 12820 1 4840							PFDPATP\LLGIYPKDYKSCCYK
12743 43111 A 12821 578 630 EYKNR*GLSLNPWGKTVHISW DVWSVGSSKAKRNWESWES*GRIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							DICT/RVCVPAALFTIANTWNPT
DVWSVGSSKAKRNWESWES*6 RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH	12742	43110	A	12820	1	4840	
RGIQKKASLRSRTGNHSASSGI ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH	12743	43111	Α	12821	578	630	EYKNR*GLSLNPWGKTVHISW
ESRV*GLGTTGYRGITASLMSL RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							DVWSVGSSKAKRNWESWES*C
RSCTRLH 12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							RGIQKKASLRSRTGNHSASSGI*
12744 43112 A 12822 8 214 EEWPKRKSEIRERPRP*SWPSD QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							ESRV*GLGTTGYRGITASLMSL
QTLVVQRGQKMEQANHPDPTI HMSQLMWTACPKGLGIALICL VRH							L
HMSQLMWTACPKGLGIALICL VRH	12744	43112	Α	12822	8	214	EEWPKRKSEIRERPRP*SWPSDK
VRH							QTLVVQRGQKMEQANHPDPTD
							HMSQLMWTACPKGLGIALICL
12745 43113 A 12823 33 240 GGGRNTWLQAVPGFPGW*RTI							
	12745	43113	Α	12823	33	240	GGGRNTWLQAVPGFPGW*RTL
							RTAVWDRRLTLRKPHHCPGGS
				ļ			QFPGPQRQVIGLLHFLSSLNHQ
GWFV						<u> </u>	GWFV

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12746	43114	A	12824	1	411	MSTITQRKEENPSVFLEWLWEA
				1		/CKKIYSPVTRLTRGSIDPKR*V
]						YYAISHRYQQKAPKPSPGP*TK
		l				SGGIIKPGNLGVLY*GPRGTGQ
						KGKARSEKGCSLSHGPQTNKP
						WWFREDRKWSRPITWYGFLSV
						WFARTP
12747	43115	Α	12825	1	611	
12748	43116	Α	12826	120	493	KYRAQRRMIARWLPQDLERGE
:		l				AMLPNSSGWYKSLKGI/VLISEA
						FSCKCRAASLTIADWLV*KQHS
				ļ		SPKKVQSPPFSAVRRSKPWQFW
						RVTAAKQSIWDCRVRIDFVISC
						KLRNPLRVCGSRILK
12749	43117	Α	12827	1	2195	VGEEGPAGVEGLNFPAQSRRLP
						GPQQCSPGAEQGASARARRPRP
						RPRPAAMVPGVPGAVLTLCLW
		i				LAASSG\SWRPAPARL\CAAAG
					RVAVCRERPARS\CASRCLSLQI	
					TRISAFFQHFQNNGSLVWCQNH	
		ļ.				KQCSKCLEPCKESGDLRKHQC
						QSFCEPLFPKKSYECLTSCEFLK
						YILLVKQGDCPAPEKASGFAAA
						CVESCEVDNECSGVKKCCSNG
						CGHTCQVPKTLYKGVPLKPRK
	Ì					ELRFTELQSGQLEVKWSSKFNI
			1			SIEPVIYVVQRRWNYGIHPSED
						DATHWQTVAQTTDERVQLTDI
l	İ					RPSRWYQFRVAAVNVHGTRGF
		1				TAPSKHFRSSKDPSAPPAPANLR
		1				LANSTVNSDG/AV*TVTIVW\DL
						PEEPDIPVHHYKVFWSWMVSS
						KSLVPTKKKRRKTTDGFQNSVI
						LEKLQPDCDYVVELQAITYWG
						QTRLKSAKVSLHFTSTHATNNK
						EQLVK\TRKGGIQTQLPFQRRRP
		1				TRPLEVGAPFYQDGQLQVKVY
			E			WKKTEDPTVNRYHVRWFPEAC
						AHNRTTGSEASSGMTHENYIIL
						QDLSFSCKYKVTVQPIRPKSHS
						KAEAVFFTTPPCSALKGKSHKPI
						GCLGEAGHVLSKVLAKPENLS
						ASFIVQDVNITGHFSWKMAKA
İ						NLYQPMTGFQVTWAEVTTESR
						QNSLPNSIISQSQILPSDHYVLTV
		<u> </u>				PNLRPSTLYRLEVQVLTPGGEG
12750	43118	A	12828	1	1410	
12751	43119	A	12829	2	299	
12752	43120	В	12830	81	510	
12753	43121	Α	12831	3	436	
12754	43122	Α	12832	1	769	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12755	43123	A	12833	15	413	ERMCAGPWRKGAETGPYFCND
						TVRHRKASLLCPF*RMNGDQN
						SDVYAQEKQDFVQHFSQIVRVL
						TEDEMGHPEIGDAIARLKEVLE
						YNAIGGKYNRGLTVVVAFREL
1						VEPRKQDADSLQRAWTVGWC
]						VELLQA
12756	43124	Α	12834	1	795	MNGDQKSDVYAQEKQDFFQHF
						SQIVGVLTEDEMGHSEIGDAIA
					1	RLKDVLEYNAIGGKYHRGLTV
						LVAFQELDAINYAILLEACIYYL
					:	LKLYCQAQPYYLNLIELFLQISC
						QPEIGQTLDLITAP/QDNVDFGR
						FT/EKQHANA*KILLEI/GEFFQI
						QDGYLDLFGDPTVTRKVGTDIQ
						DNKCSWLVVQCLQRSTLEQYQ
						ILKENYGQKKAKKVVQVKALY
						EELDLPAVFLQYEEDGYSHIMG
						-
10757	42125	 	12025	1	328	LTEQYAAPLLPAMFLGLVYKIY
12757 12758	43125 43126	В	12835 12836	2	868	
12759	43126	A B	12837	118	1138	
12760	43127	A	12838	100	374	KHISPALKALELPFERNLII/PLQ
12/60	43128	A	12030	100	374	LLKVRILKEGEML/DI*HWFL**
						4
						GMGTVQKGMPHKCYHGKTGR
12761	43129	A	12839	1	1035	VYNVTQHAVGIVVNKQVK*VV
12762	43130	A	12840	13	600	INPPPPPFRPELPSSSNSPKMTDH
12702	1,5150	'	120,0			KGERGEATRYMFSRPFKKTMG
						VVPLGHNN*RF\YKKGDIVD\IK
						GMGYCVQNGMP\HKCYPWPKL
						EGVLQLLPQHA\VAIVVNQPVL
						GQSFFPRE*IVRIEHIKHF*EPEIS
						FLK/RVLKENDSEKERSPNEKG
		ŀ				TWGSNLKRHLAPPQKKHTL*R
		İ				TNGKEPEL\LEPIPYEFHGHN
12763	43131	A	12841	1	317	QRPSEAKEIKLYAQIPPIEKMDA
12/03	ונונד	^	12041	'	317	SLSMLANCEKLSLSTNCIEKIAN
-						LNGL\EAVGDTLEELWISYNFIE
					1	KLKGIHIMKKLKILYMSNNLVK
}		1				DWGTPVIKGDEEEDN
12764	43132	A	12842	155	588	D # O I I A IZODEREDIA
12765	43133	A	12843	1	954	
12766	43134	A	12844	1	486	
12767	43135	A	12845	1	367	-
12768	43136	A	12846	1	190	LISRAKEDDMTLNIAEGVHPSY
,50			1	_	-	NIVLNTLHT/CGISVKSTWRPSK
						GAAP*ARMPGGTAGPCLAS
L		Щ	L	<u> </u>	L	onn man como cons

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12769	43137	A	12847	1	2376	MRGAQKATNVNKLSEDIQGKE
1.2,00	1.5.15.			-		ESPTQFYERLWEAYRMYTPFDP
			:			DSPENQRMIPMALVRQSAEDM
1						RRKLQKQAGLAGMNPSQLLEI
						ASQVFVNRDAVSRKENGKENG
						GQARRYADLFSRTKNYQPVQD
						LRLLHQAKLTFHPTVPNPSTLL
						GFPPAEDSWFTCLDLKDAFFPIR
						LAPERQKLFAFQWEDPESGWPP
						CWRA\WQQLPC*YKIVPSILRIS
						SSDGKYKAFSTCGSHLAVVC
12770	43138	A	12848	1	636	MRGAQKATNVNKLSEDIQGKE
12//0	43130	^	12040	1	030	ESPTQFYERLWEAYRMYTPFDP
						DSPENQRMIPMALVRQSAEDM
						RRKLQKQAGLAGMNPSQLLEI
						ASQVFVNRDAVSRKENGKENG
						GQARRYADLFSRTKNYQPVQD
						LRLLHQAKLTFHPTVPNPSTLL
						GFPPAEDSWFTCLDLKDAFFPIR
						LAPERQKLFAFQWEDPESGWPP
10771	42120		10040	100	011	CWRA\WQQLPC*YKKQIS
12771 12772	43139	B A	12849 12850	180	811 1265	FLGVPTLGGWRPIQHELRK/PRE
12//2	43140	A	12030	1008	1203	VTPVLLRDLIRRFRPPLWIGSDN
						GPAFLAALVQKTAKGIQNNITG
						GVYTLCDIDSHIILFRSGY
12773	43141	В	12851	1	996	OV I ILCOIDSHIILF RSG I
12774	43142	A	12852	1251	1424	DSPRGEAES*A*LPEKLLEMSG
12//4	43142	l'A	12032	1231	1424	NRLVDNKDPGKKQTQRRIPHSP
						SQIPVPLPEIWCT
12775	43143	Α	12853	1	1121	SQII VI EI EI WCI
12776	43144	A	12854	189	621	
12777	43145	A	12855	2207	2308	
12778	43146	A	12856	1	762	
12779	43147	A	12857	1	678	
12779	43148	A	12858	3	496	
12781	43149	A	12859	107	258	NALGKRTSFTNRNFLASGRLFD
12/81	7,777	<u>ا</u> ``	12000	10,	250	SVTYARRLQQYPAPL*KTAI*SF
						SLLNH
12782	43150	A	12860	2	410	SPDPWVSTYKSTCHIAQEIAEKI
12/82	43130	/A	12000	 	410	· .
						HLRNQYERKGE*AP\RSNLMSE
						DANGGAPNPWLFEEPEETIGW
						GFDEIRQQQHNIIR*QDAGLDA
						LSSIISRQKQM\GIEIGNELHEQN
						EIIDDLANLMENTDEKLRNETR
12792	42151	A	12861	3	1402	RVN
12783	43151	Α	12001	2	1402	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
12784	43152	A	12862		711	MAPDPWFSTYDSTCQIAQEIAE KIQQRNQYERKGEKAPKLTVTI RALLQNLKEKIALLKDLLLRAV STHQITQLEGDRRQNLLDDLVT RERLLLASFKNEGAEPDLIRSSL MSEEAKRGAPNPWLFEEPEETR GLGF\DEIRQQPQKII\QEQDARL DALSSIISSQKQ\MGQEIGNELD EQNEIIDDLANLVENTDEKLRN ETRRVNMVDRKSASCGMIMVI LLLLVAIVVVAVWPTN
12785	43153	A	12863	1	936	
12786	43154	A	12864	1	1023	
12787	43155	A	12865	1606	2238	
12788	43156	A	12866	3	347	
12789	43157	A	12867	1	1665	
12790	43158	A	12868	1	589	
12791	43159	A	12869	19	1677	
12792	43160	Α	12870	1	906	
12793	43161	A	12871	1	228	
12794	43162	Α	12872	1	320	
12795	43163	A	12873	1	398	TSKAPGAQGEQGFEECFLAVAL AGRPLPSLWALFQATTPV\LQG LKEAFFRPEVP\LRRDLPLLLFR\ TQTSDPAMLP\TMIGLLAEARR AGCLSYQTSLVSDGETWHVMG ISSLLP\SLEG\WKQ\HVTEAFQF HF
12796	43164	A	12874	3	322	SAGGSGRRTLHSRTMAQFVRN LVEKTPALVNAAVTYSKPRLAT FWYYAKVELVPPTPAEIPRAIQS LKKIVNSAQTGSFKQLTVKVTT G*MKTDVHNRKCLPLGFF
12797	43165	A	12875	91	425	WTFHPTTMAPFVRDLGEKTPA LGKAAVTYLKPRLAAFWYYH\ QVELVPPTPA\EIPRAIQSLKKIV\ NSAQTGSFKQLTVKEA\LLNDL VATEVSTWFYVREITGKRGIIG* NV
12798	43166	A	12876	179	664	HSSPAVPGRTDFSHSQNPWPQF CPVTLVGEDPRALVNAAVT/YT RKPRIGLHFWYYA\KVELVP\PT P\AEIP\RAIQSLKK\IVNSAQTGT LANSSHVKEAV/L*NGLVATEV\ LMWFLCSERLLGKRGHSLGYG CFEDQSFNIWIIFWILFILELFVW DHVVDPDC
12799	43167	Α	12877	25	164	
12800	43168	A	12878	2110	2260	
12801	43169	C	12879	32	1429	
12802	43170	A	12880	172	322	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12803	43171	Α	12881	1060	12039	
12804	43172	Α	12882	1	810	
12805	43173	Α	12883	3	146	
12806	43174	Α	12884	1	1211	GPQDGNQQPAPPEKVGWVRKF
						CGKGIFREIWKNRYVVLKGDQ
						LYISEKEVKDEKNIQEVFDLSD
						YEKCEELRKSKSRSKKNHSKFT
					}	LAHSKQPGNTAPNLIFLAVSPEE
						KESWINALNSAITRAKNRILDE
		ŀ				VTVEEDSYLAHPTRDRAKIQHS
						RRPPTRGHLMAVASTSTSDGM
		ĺ				LTLDLIQEEDPSPEEPTSCAESFR
						VDLDKSVAQLAGSRRRADSDRI
		!				QPSADRASSLSRPWEKTDKGAT
						YTPQAPKKLTPTEKGRCASLEEI
		ŀ				LSQRYAASAHTLQLRAEEPPTP
		ŀ				ALPNPGQLSRIQ\DLVARKLEET
						QELLAEVQGLGDGKRKAKDPP
						S/RSPPDSES\EQLLL\ETER\LLG
						EASSNWSQAKRVLQQVRELRD
						LYRQ\MDLQTPD\SHL\RQTTPH
		ļ <u> </u>				S\Q\YRKSLM
12807	43175	A	12885	617	1041	TTLINPGGPAPLLSLPTSSITGFT
						TFYHPNLKPLRTCPS*VQAWPC
						RHPI*SRDYQRDAEPTTCISKLL
						GENIRRDEFWKVQIKARSGETE
						QNPRSTAFAFGELPLANREELN
		}				LLLNKGILPLCDQRQEIESECAF
						TIFCAGDANCFPPSPTHLPLTAL
						QQRPCGLSGSASKEAWGNSLV
						CGPRGDTLPMAIAKPHRSASKH
						STLHQYPTPPRSLCHQHRDEQP
						TNHLQTLNTIPAWIHTKWVPGL
						DHINKPWRSCSSSQSAYLFYHW
						LHYFLPPKLKTSKDLSILSAGLA MSAPHLIFRVSAITSNSSCKPSS
						TSKACEELFKEHQCLGSTDIPIH
						IQGGAPFSAVSPQRLVLRGSQD
Ì						GSHMVRPRLPFPDSAGLANFHL
						FSKLFLD
12808	43176	A	12886	1	606	I SKULDD
12809	43170	A	12887	174	350	VSWRFVQVFI*V\AGGSRICAAA
12009	77177	^	1200/	'/7	UCC	GLSPGTPPPCATSLIGACSLIFIP
!		1				RLLSLFCGDEAL
12810	43178	Α	12888	3	265	TEDODI CODEAL
12811	43179	A	12889	1	1209	
12812	43180	A	12890	53	369	VAHQEKGIAEPGLRPTGDSRQT
1-012		•	,			HRRLDVERSTSVQEHMGGCHF
						SPFPEREKLSTLRGIHQQAPAL
						WQATDQWIDIEF/GLGQPEESL
						GR*ITRLQGKTILPFGSPIC

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12813	43181	Α	12891	1	345	
12814	43182	A	12892	2	451	
12815	43183	Α	12893	2	286	
12816	43184	A	12894	217	1071	GENPRSENTRLATILEVACCHF GSSPPPS\GSSGRKDPPLTFGDH EGTSKAEPSPALGLTPEHKGNV GHAARIQQPGPLSLWSRKAGK GVQDCYIDLKQIKIDLDKFSDN PDGYIDVLQGLGQSFDLTWRNI MLLLNQTLTPNERSATITAARE FGDLWYLSQVNDRIAAEKREK FLTGQQAVPSVGSHWDTESEH GDWCCRHLLTCMLEGLRKTRK KPMNYSMMSTITQGKEENPTA
12817	43185	A	12895	1	756	FLERLREALRKHTSLSLDSIEGQ LILKDTFITQSAADIRK\NFKSPP
12818	43186	Α	12896	473	1015	
12819	43187	С	12897	152	322	
12820	43188	A	12898	146	1079	PPCAVVCLLWIAAVYCLSTTFC APLCGQNTWLPKPCRHVNRAA SNASVSLSTVTLFSRLS*PQRKG SELLDSSGPLPASPSLPLCSGVS PRAGLGSAWPKIPFLGIREAKN PRSENTRLATILEAGHRHLGTS VSKDHPVTFWRPRRDLQSDLK QIKIDLGKFSDNPDGYIDVLQEL GQSFDLTWRDIMLLLNQTLTPN ERSATITAAREFGDLWYISQVV AAVAGLVSEAVKIIQGSYCVDI HDVNGILTAKGDLWLSDNHLL KYQALLLEGPVLRLRTCATLNP ATFLPDNEEKIEHNCQQVIAQT
12821	43189	Α	12899	189	1798	

SEQ ID	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
12822	43190	A	12900	194	1729	NPAAQTPFFVVKKGGKRGAGL
						LHRQYPLRLEAKQGLKKIVKDL
						KAQGLVTPCSSPCNTPTLAVQK
						PNGQWRLVQDLRIINEAVVPLY
						PAVPNPYILLSQIPEEAEWFTVL
						DLKDAFFCIPVHPDSQFLFAFED
						PSNPMSQLTWTVLPQGFRDSLH
						LFGQALAQDLSQFSYLDTLVLQ
						YMDDLLLVTHSETWCHQATQA
					!	LLNFLATCGYKVSKQNAQLCS
						QQVKNLGLKLSKGTRALSEERI
		İ				QPILAYPHPKTLKQLRGFLGITS
						FCRIWIPRIKAVKLQIILQMEPQ
						MQSMTKIYHGPLDRPTSPSSNV
		ļ				NDIEGTNASDFPFLSQVLEQVV
						SPKGSKEAQCCVLRPLGCESNL
						KQIKIDLGKFSDTPDGYIDVLQ
						GLGQPYYLTWRDIMLLLDQTL
		l				TPNERSAAITAVREFGDLWYLS
		ĺ				QVNDRKITEEREQFPTGQQAVP
						SVDPHWDTESEHGDWCHRYLL
						TDVLEGLRKTRKKP\IN*SMISTI
	<u> </u>	ـــــ				TQGKEENPTAFLERLREALRKH
12823	43191	A	12901	54	254	
12824	43192 43193	C A	12902 12903	32	221	NGVGHNRMGTNGSGGGEWG\P
12023	43173	A .	12903	32	221	PAPASA*GCSLASAFASASPRW
						WFEMGCRGIYDAQQVALSRSF
12826	43194	A	12904	203	538	WI EMGERICATION OF THE PROPERTY OF THE PROPERT
12827	43195	В	12905	131	536	
12828	43196	A	12906	1	1251	
12829	43197	Α	12907	38	475	ESERRGEFCLCKIISEQGRAVCV
						EDEQTSHFALTGNNGIYRKSQQ
		1				QWQQELSDLARDPPAQCSAGP
						VGDDMFHWQATIMGPNDSPYQ
						GGVFFLTIHFPTDYPFKPPKVAF
		1				\TTRIYHPNINSNG\SICLDILRSQ
						WSPAFNQFPKSS
12830	43198	Α	12908	282	849	QSCPAAGIIKSFGRTFEHTPREV
						RKPDDKHTLLALKRINKDLSDL
						ARDPPAQCS\AGPV\GDDMFHW
						QATIMG\PNDSPYQGGVFFLTIH
		1				F\PT\DYPFKPP*GLHFTTRFY\HP
		1				\NINSKWAAFCLRYS*DSQW\SP
		1				ALTFSKVLLSI\CSLL\CDPNP\D\
						DPLVPGDWHGSY*NRTGDKYN
						RISREWTQKYAM
12831	43199	A	12909	1	397	
12832	43200	A	12910	245	535	
12833	43201 43202	В	12911	265	333 562	
12834	1+3202	Α	12912	1203	1302	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
:	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
12835	43203	A	12913	11	872	MVNTECQLDWIEGCKVLILGVS
						KEASQGLGGRSRGRNVRQSTD
						ADIRPTSEKDARSLMTTRQEDN
						ISVDQRGFLSGVQKKSGCANGL
						KGSVCGGFYWVMAVIRMGVG
						KAMGLNLKLGIKFEIKKGYLKG
						CRDSLRLSPTGFGKFTATSHEV
						AATVPIKVVLELTLQTHKILRG
						VFAHLTSCLQLHSLPPGSSMDS
						AKARRQLQVAGQGHSLVKQK
						GFHVEAPDHHPVQLSDISCKSY
						RWGHKLIKINFLIPPLWWYYDL
						EEVKREQSEEKQQAKIMADIIG
						DALNAQKAFKGNPKGHKPLPF
		1	İ			SAGQMENW*CPECPKSI
12836	43204	A	12914	7	231	SAGGMENT OF ECTRO
12837	43205	A	12915	1	223	MEILTTCDKGLIFTGNTRKSETS
						TSECTCPLDINCSLSHSPPSPSDE
					1	ETQHHTDSLHVRPRKGYLLCIY
						HIPDIVAVMVNIQCQLDWIEGC
						KVLFLGVSVKVLPKEINTGISGL
						GKADPPSIWKS*QHVTKDLYLQ
						VILENQKLVPANALVRWILTVP
		1				FPTLLRALQMRKHSTTLTVCMS
						DRGKDIYCASTIYQI
12838	43206	A	12916	1083	1273	NKMSFKLKEDCFGRSKCNL*T*
						CWKSINEA*K/DLELQLQFGPQE
						KLFALEQQVKKKLNLVPDA
12839	43207	С	12917	1	1506	
12840	43208	Α	12918	3	1219	
12841	43209	Α	12919	1	735	
12842	43210	Α	12920	2	94	
12843	43211	A	12921	362	460	WLILSVNLIGLKDAKYCSCVCL *GCCQRRLTF
12844	43212	A	12922	1235	1474	
12845	43213	Α	12923	1	250	
12846	43214	A	12924	1	261	
12847	43215	A	12925	2	85	
12848	43216	Α	12926	1	291	
12849	43217	A	12927	1	321	
12850	43218	Α	12928	1	370	
12851	43219	A	12929	1	684	
12852	43220	С	12930	1	210	
12853	43221	A	12931	2	91	
12854	43222	A	12932	1	243 353	WMVI ETIII CVI S*COVTVIIDA
12855	43223	A	12933	2	333	WMKLETIILSKLS*GQKTKHRM SSLIGSTDQQRLVVKNVQGRVC
		1				PGSTGHGAWEAWTAMRKVDS
						PLQTGSLRPGWCRQVGSHARS
1						GSAEAAGNPGLHPGICPRVEDG
						IQVAENFPEV
<u> </u>	<u> </u>	1	L	<u></u>	L	IQVAENTEV

1 -	SEQ ID NO:		1 -			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12856	43224	I C	12934	1	289	
12857	43225	Α	12935	1	1194	
12858	43226	Α	12936	140	237	
12859	43227	Α	12937	1	292	
12860	43228	Α	12938	2	178	
12861	43229	В	12939	1	1245	
12862	43230	Α	12940	2	241	
12863	43231	Α	12941	1	675	
12864	43232	Α	12942	1515	1920	TPTNLQEKNKQPHQKVGEGYE
						QTLLKRRHLCSQKTHGKNAHH HWPSEKCKSKPQ\WIPSISHQLE WQSLKSQETTGTWMKLETIILS KLLQGQKTKHHMFSLIGGNRT RRTHGHRKGNITLWGLLSDSV NCASSSG
12865	43233	A	12943	608	891	THE RESERVE
12866	43234	$\frac{\Lambda}{\Lambda}$	12944	1	3956	MGSCPGGFTAEFYQRYKEELVP
						FLLKLFQSIEKQGTLPNSFYEAS IILIPKPGRDTTKKENFRPISLMN IDAKILNKILTNRIQQHIKKLIHH DQVGFIHGMQGWFNIRKSINVI QHINRTNDKKHMIISIDAEKAF DKIQQPFMLKTLNKLGTWMKL ETIILSKRSQGQKTKHRMFSLIV NIIPTIKILIKQNMTAMETLCPIF KLEGDTMDNPIANRRVRTAVP TSVVCLHVGTEPEARLPHGPMI TAQSPFG
12867	43235	Α	12945	1	855	
12868	43236	Α	12946	1	255	
12869	43237	Α	12947	273	3484	
12870	43238	Α	12948	3	377	
12871	43239	Α	12949	85	333	
12872	43240	Α	12952	22	308	
12873	43241	В	12953	30	383	
12874	43242	Α	12954	3	448	
12875	43243	Α	12955	1	453	
12876	43244	Α	12956	3	491	
12877	43245	Α	12957	2	749	
12878	43246	Α	12958	155	637	
12879	43247	Α	12959	1	510	lanus and an an an an an an an an an an an an an
12880	43248	A	12960	2	559	RPHASAHASGRQRSQDVTTMV WALLLTLLTQGTGSWAQSAL TQPPSASGSPGQSITISCTGLIND VASYNFVSWYQQHPGKAPKL MIYEVTKRPSG\VPDRFSGSKSG \NTASLDHLWGSRLEDEADYYC LLICRRLHPGGGIRRWDPRVTV LGSAQRPTPTVTSGSRPSSEGAP SQQGHTSVSDQ

SEQ 1D	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12881	43249	Α	12961	1	1958	MSVQRKDRVRTQQGGSVASSM
ļ						LTQEPVGSVALGQSARDRDPD
						CSDSVSVYLMDTWQSVIIGFAE
ĺ						SAATLGDPPELPMHPEGSEASY
						ELTQPPSVSVSPGQTARITCSGD
		ŀ				ALPKQYAYWYQQKPGQAPVL
		ŀ				VIYKDSERPSGIPERFSGSSSGTT
						VTLTISGVQAEDEADYYCQSAD
						SSGTYPTVTQADRETPTHQYSR
						HPSYAKGFVFLWGSWAQSALT
		1				QPASVSGSPGQSITISCTGTSSD
					ļ	VGSYNLVSWYQQ\PQGKAPKL
						MIYE/VDSKRPSGVSNRFSGSKS
						G\NTASLTISGVPGLRDEG\DYY
					!	CCSYAGSVASYELTQLPSVSVS
						PGQTARITCSGDVLGENYADW
						YQQKPGQAPELVIYEDSERYPG
		l				IPERFSGSTSGNTTTLTISRVLTE
				:		DEADYYCLSGDEDNPSVTQAD
						GEDTGIRSHYEGTLALYLYPEP
						VKATCSVTSYVLTQPPSVSVAP
	1	l				GQTARITCGGNNIGSKSVHWY
		l				QQKPGQAPVLVVYDDSDRPSGI
		l				PERFSGSNSGNTATLTISRVEAG
						DEADYYCQSLSTLGPKTLGDTV
		l				LRPMIYSVSIQASGVPDGFSGSK
		l				SGNTASMTISGFQAEDEADYYC
						NSHRRGSVVSSELTQDPAVSVA
						LGQTVRITCQGDSLRSYYASW
						YQQHPGKAPKLIIYDGH
12882	43250	Α	12962	2	367	
12883	43251	A	12963	2	376	QTYSLRRATPRHIIVGFTKVEM
						KEKVLRAA/NKPIRLTVDLSAET
						LQARKEGGPIFNILKEKNFQPRI
						SYPAKLSFISEGEIKSFTDKQML
			1			KDFVTTRPALQELLKEALNME
						RNNQYQPLQKHAKW
12884	43252	A	12964	2	1178	
12885	43253	Α	12965	1	732	
12886	43254	В	12966	99	716	
12887	43255	Α	12967	1	1011	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12888	43256	A	12968	3	1263	GRTIQTKGKEV\ENFEKNLEECI
12000	43230	^	12906	5	1203	TRITNTEKCLKELMELKTKARE
		ŀ				VREECRSLRSQCDQLEERVSAM
						EDEMNEMKQEGKFREKRIKRN
						EQSLQEIWDYVKRPNLRLIGVP
						ESDVENGTKLENTLQDIIQENFP
						NLARQANVQIQERQRTPQRYSS
						RRATPRHIIVRFTKVEMKEKMG
						LLVPNWTNHSPLFRAILFDYKG
						FCRFGTTHQTGFSPAGANQRGP
						LAATLSGPGGEGQSAVARLTGL
1						IGVPESDVEKETKLETTLQDIIQ
						ENFPILARQANVQIQEIQRTPQR
į						YSSRRATPRHIIVRFTKVEMKE
						KMLRAAREKGRVTLKGKPIRL
						TADLSAETLQARREWGPIFNIL
		1				KEKNFQPRISYPAKLSFISEGEIK
		1				SFTDKQMLRDFVTTRPALKELL
						KEALNMERNNRYQPLQNHAK
12889	43257	A	12969	1	1392	nts. is in its i
12890	43258	A	12970	1	1224	
12891	43259	Α	12971	1	970	
12892	43260	Α	12972	1	954	MDGDLHKGVAVFWPPDAAGG
						HTCCRSMVCRSLLGGAQWKFH
						SFRCRIRFELILHMIQERNSINIS
ļ			ļ.			QKDVHTETPSETHHHQRPKVD
ŀ						KSMKMRRNQCKKAENSKNQK
					1	ASSPPKEHNSWRAREQNWTEN
}						EFDKLTEVGFRRWVITNSSELK
						EHILTQCKEAKNLEKTLEELLT
						RITSLEKNINDLMKLKNTA*EL
						REAYTSINSQISQAEERISEIEDQ
						LNEMKHEHRIREKRMKRNKQS
						FQEIWDYVKRPNLRLISVPESD
						GENGNK*ENILQDII\QENFPNL
						ARQANIQIQEIQRTPQRYSSRRA
						TPRHIIVRFTKV
12893	43261	В	12973	1	1346	
12894	43262	Α	12974	1	318	GSRGELLLSLCYNPSANSIIVNII
						KARNLKAMDIGGTSDPYVKVW
						LMYKDKRVEKKKTVTMKRNL
						NPNFNES\FAFDIP\TEK\LRETTII
						ITVMDKDKLSRNDVIGK
12895	43263	Α	12975	17	389	SWGPHCTHKLKKSFSLMPEMT
				[DAMDRERM/RGSKKDV/R*VLG
						GCAATWTPRAGGRQQSPGTTV
						DKREDTPDCAMCDQSTSHPISA
						SSPSLLPGSSFTTQTFPPGIAHYR
<u></u>		<u></u>				ASFLIVAYYPSNKNSK

SEQ ID			SEQ ID NO		Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
12896	43264	A	12976	84	282	GEGQVGWPQGEVQQDEEVIEE VGA*DAGSGYQEEEEGPQSLRR EVVGVELVVGADPSGEGPKED
12897	43265	Α	12977	1	1179	
12898	43266	Α	12978	3	634	
12899	43267	A	12979	234	619	CYVVQEVLPSLLCHCLVICSRP GRTAEWPPG*GF*SPPHRGWW GC*GHD/HRRGTSIFPTH/V*SPP ASYPGSLCRPGGQEEREREGEW QREGRRPGTR*CGAEAEARGW RAAAKATAVAGARNRRPGISG
12900	43268	Α	12980	1	1182	
12901	43269	Α	12981	3	763	
12902	43270	Α	12982	32	949	_
12903	43271	A	12983	1	1299	
12904	43272	A	12984	1	1176	
12905	43273	Α	12985	1	421	
12906	43274	A	12986	81	404	VKFGPEIWCRDSDQGRGGVGT SLGRSIPCPPALCSVRKIYLRPL VLRPTSPRNISPILNRDPTVQLT WQPLPEPLELWPKAL*LTPSQIY SA*RLKTDAARLPRKPPR
12907	43275	Α	12987	1	378	
12908	43276	С	12988	1	751	
12909	43277	A	12989	302	629	ICLSYPAPPCKRFPHYTRVLYH* SLFNENTSQSCFTSR/WKLESFT TRAIERHQIPLL*ATLTLIRWLD KKLAFQLLSLTASFSPSYQSLPP TPPLKLPPIISSHTRQMV
12910	43278	Α	12990	14	380	
12911	43279	A	12991	189	608	QSDLSASQHGLFPLATEVRSSG AASCPDSDSICPAPTIAPGRATP PQANCWASEGTLRYQALPGNR APVSQVSQAGGANLCVQQPKK HLTNFKSGKRPLFTLFSNLQGP RSRPVAFLSKQLDLTVLGWPSC L\VQQLLPP
12912	43280	Α	12992	1	513	
12913	43281	A	12993	2	462	WSMGLPRRSGCVSLQPLSQEDL GRSQSESLGPEFQGLWEWLPEQ LPRSFIQALGSLSYFPRTLMNLS WPLCLRKWGYL/SLYLPFQVGL PSTVPWPLILPSKVPEEEGQLQS TLPCGIKLLPVFTSHHAFLGVFC NQWVSDVFLALFLMKLLGAQ
12914	43282	Α	12994	59	213	
12915	43283	Α	12995	1	1159	
12916	43284	С	12996	1	864	
12917	43285	Α	12997	1	1155	

Unknown,
nucleotide tide insertion)
CKMSGRGP
GTVLWLW
FLQKLPRL
LWNCLFLL
PLTT*HS*P
ARAVTPFV
CSLCLDASV
RLRWSFGL
RRK
KKK
SYSGYWPE
SEVLYSALR
KPRRSGKE
GVGVREID
RLLQSHYGI
AALYQLNV
SILDSVQPN
GARNLRA
MENVSALS
DDSLERVK
FLGLAGNG
VNTCLPSDP
VLEQLHTS
NLFLCSLLD
YNQYDLQ
IPW*PSAFV
VGGGDDN
GDRAVSVF
AL/VPEFQG
GIDTILCHH
SLMSVGNS
ADTVGLII
SNKHIELRT
MTSSYNQE
VESPTSMF
DIFSMIIYT
WVVGFHM
HGPLTCDP
SNSGALED
QGNSWWK
TQGSHQL
TAHFFFGFF

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12926	43294	A	13006	3	371	AGARFNVRSH*RNDRVRPHRD VYSLQGRLSDHSPTFQGCQTTQ GRLPWSFTLSGKFRFSGEGATT SPAHKNFQTPEPQWPGIPPEPPP TGACYTCRKSGHWAKECPQAR IPPKPHPICVGPH
12927	43295	A	13007	77	332	RWKKNCLSPPRMS*QR*WKP* MQQAACSVGETQPPCTQVK*T ALLLTQSLFGGLFTRTHMKFGA VTRIGGPPLGDQSPVLLLFAP
12928	43296	A	13008	1048	1349	RVVEQDDAERLFRRFRAA/GDL PYGAFGEIF*ETQCQEMQQLAE DHASVTMISLEAWRLYRKMVC RIWASMLRLSVCHLCAPKRVW SLISRNGYHLPGSIH
12929	43297	A	13009	213	499	PEQRTSHRQQQRRQPPPQSRS*P FCPSLPRSALSLSFLCDGYAAP* AYRE*IPVTEDHFAE*I/ISWCPC *YREALGQLLAKMRR*SARKRF QLSP
12930	43298	Α	13010	2	431	
12931	43299	Α	13011	2	367	
12932	43300	В	13012	1107	2083	
12933	43301	В	13013	1	627	
12934	43302	A	13014	712	2671	AHVGRGST*ALPRAMGSDAF*C PLQ*MTPASFGSKAALSRVFIKE ALMMEDPCVVRKPLSAHMTA WWCRIWKAYLESV
12935	43303	A	13015	50	1362	WEQIPHGLVLSLQQLPRRSGSIS LQPLSQEDLGRSQSESLGPEFQ GLWQWLSDFQT*HQWFVSGFQ AFSDRLKAALSASLLLRFGDSD WLPSSSACKCLMLGLHFVIVGN ICATLKEKYSSMLHLDVTMKK NGEKRTRLQKRKKGMPPHPAY EDLNIAAITLPANVVLHQPSGFR TSGQLDPVWWSLDTDAHEIWC QDPGLGSGDFPWEITPLSSYSLL HEKDPPTTSGPQTDQPKKHLTN FKSKTKETGFIHGPKTPAPVTD WEGSLPLVFNHCRDTSLIIHPCF KGVRPRRDACLGPSPLAASPAF LEKGQDLINLAFKVYNNRKKL QFLASTVRQTPATSPAHKNFQT TEPQQPGVPPEPPPTGACYMCR KIWPLGQANARSPGFLLSRVPS VWDPTGNRTVQLTWQPLPEPL ELWPKALCLTDSFPDLLGLAA
12936	43304	A	13016	1	507	
12937	43305	Α	13017	5	271	
12938	43306	A	13018	1	618	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12939	43307	A	13019	1	429	
12940	43308	Α	13020	1	3567	
12941	43309	Α	13021	1	353	
12942	43310	A	13022	1	338	
12943	43311	Α	13023	1	1175	
12944	43312	Α	13024	3	326	
12945	43313	A	13025	150	494	PSPSLGYLVGTRGTALRL*DAR AAMRPFDPSTLLPTCWDYWTY AGSLTTPPLTESVTWIIQKEPVE VAPSQLSAFRTLLFSALGEEEK MMVNNYRPLQPLMNRKVWAS FQATNEG
12946	43314	A	13026	2	975	DSREHHPLQTGVSGNRPKMLG RNTWKTSAFSFLVEQMWAPLW SRSMRPGRWCSQRSCAWQTSN NTLHPLWTVPVSVPGGTRQSPI NIQWRDSVYDPQLKPLRVSYE AASCLYIWNTGYLFQVEFDDA TEASGISGGPLENHYRLKQFHF HWGAVNEGGSEHTVDGHAYP AELHLVHWNSVKYQNYKEAV VGENGLAVIGVFLKLGAHHQT LQRLVDILPEIKHK\DARAAMR PFDPSTLLPTCWDYWTYAGSLT TPPLTESVTWIIQKEPVEVAPSQ LSAFRTLLFSALGEEEKMMVN NYRPLQPLMNRKVWASFQATN
12947	43315	Α	13027	1	1233	
12948	43316	A	13028	761	1000	IPFISFSCLIALARTSNTMLNKSG ERGHPSLVPVFIGNASSFCPVSM ILAVGLS*IALIILRYVPSIPNLLR VFSMKGC
12949	43317	В	13029	697	1527	
12950	43318	Α	13030	1	1695	
12951	43319	A	13031	434	925	VAKIFSHFVCCLFTLMVVSFAV QKLFSLIRSHLSILSFVAIDFGVL DMKSLPMPMS*MVMPRFSSRV FIVLGLTFKSLIHLELIFV*GVRK GSSFSFLHMASQFSQHHLLNRE SFPHCLFFSGFSKIR*L*ICGIISE GSVMFH*SISLFWYQYHAVLVT VAL
12952	43320	A	13032	1124	1234	KITPKKQIRQSPNLTQNTGQQ*S LQLWEHFQISTHF
12953	43321	Α	13033	929	1261	
12954	43322	A	13034	1118	1354	IPFISSCLIALARTSNTMLNRSGE RGHPCLVPVFKGNASSFFPFSMI LAVGLS*IALIILRYIPSIPSLLRV FSVKGC

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
12955	43323	A	13035	2	348	HQHLLFPDFLMIAILTGMRWYL
		ŀ				IVVLICISLRASDDEHF/YHVSV
						GCINVFF*EISVHILRPLFDGVV*
						FFLVNLFKFFVDSGYEPFVRWV
						DCKNVLPFCRLPVHSDGSFFCC
						AEAL
12956	43324	Α	13036	578	712	LISNFSKVSGYKINVPKSQAFLY
		<u> </u>				TKNRQTAKS*VNSHSQLLQRE
12957	43325	Α	13037	1	1833	
12958	43326	В	13038	1	1587	
12959	43327	В	13039	1	1851	
12960	43328	Α	13040	1349	1444	
12961	43329	Α	13041	953	1277	MVLPRFSSRVFMVLGLTFKPLI
						HLELIFV*GVNFV*GSSFSFPHM
						ASQFSQHHLLNRESFPHFLFLSG
						LSKIR*L*MCGIISEGSVLFHWS
10060	12220	 	120.40	004	1006	ESLFWYQYHAVLVTVAL
12962	43330	В	13042	804	4236	
12963	43331	A	13043	1	744 797	
12964	43332	A	13044	137	418	
12965	43333	A	13045 13046	210	927	
12966 12967	43334	A	13046	1	1415	MGFLGTGTWILVLVLPIQAFPK
12907	43333	A	13047	l'	1413	PGGSQDKSLHNRELSAERPLNE
1		ŀ				QIAEAEEDKIKKTYPPENKPGQ
						SNYSFVDNLNLLRAITEKEKIEK
						ERQSIRSSPLDNKLNVEDVDST
						KNRKLIDDYDSTKSGLDHKFQ
	ŀ					DDPDGLHQLDGTPLTAEDIVHK
	ļ					IAARIYEENDRAVFDKIVSKLLN
	-					LGLITESQAHTLEDEVAEVLQK
			İ			LISKEANNYEEDPNKPTSWTEN
						QAGKIPEKVTPMAAIQDGLAK
				:		GENDETVSNTLTLTNGLERRTK
						TYSEDNFRDFQYFPNFYALLKSI
						DSEKEAKEKETLITIMKTLIDFV
	i				}	KMMVKYGTISPEEGVSYLEGL
1						DEMIALQTKNKLEK\NATDNIS
						KLFPAPSEKSHEETDSTKEEAA
						KMEKEYGSLKDSTKDDNSNPG
						GKTDEPKGKNRSPIWKPIRK\NI
i	ŀ					EWLKKHDKKGNKEDYDPFQR*
						EVSFNNKADA\YVEKGILDKEE
			<u> </u>			AEAIKRIYSSL
12968	43336	A	13048	1	879	
12969	43337	Α	13049	3	640	
12970	43338	Α	13050	134	382	DGRQLRRGSLGGALTGRYLLP
		1				NPVAGQAWPASAETSNLVGMR
						SQALGQSAPSLTASLPCVSKCW
	1	 			1.00	ENIPPWMN/WPPH*IKHIQS
12971	43339	Α	13051	87	500	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
12972	43340	Α	13052	1	1356	
12973	43341	A	13053	1	447	
12974	43342	A	13054	3	107	
12975	43343	Α	13055	3	2377	
12976	43344	Α	13056	1	2193	
12977	43345	Α	13057	1	2724	
12978 12979 12980	43347 43348	A	13058 13059 13060	1 1	819 623	QSSIPAIMSSFLGTGAETMPVPL GAGMRRTSTEPQRVCFCTGIIF KTSSLREAPRKKSMISDSLMGR E/WMPVTKLGRLVKDMKIKSLE EIYLFSLPIKESEIIDFFLGASLKD EVLKIMPVQKQTRCGSVLVRLI PAPRGTGIVSAPVPKKLLMMA GIDDCYTSARGCTATLGNF/AK ATFDAISKTYSYLTPDLWKETV FTKSPYQEFTDHLVKTHTRVSV QRTQAPAVATT MARRKNGKWESEIIDFCLGGSL KDEVLKIMPVQKQTRAGQCTR FKAFVAIGDYNGHIGLGIKCSK E/LATAIRRAIILAKLSIVPVCRG YWGNKISKPHIVPC\RGTGIVSA PVSKKLLMMASINDCYTSARG CTAILGNFDKATFDAISKTYSYL TPNLWKETVFTKSPYQEFTDHL
12981	43349	A	13061	1	949	VKTYTRVSMQSTQAPALAKLP RVAVQPRAEV KWRITPVPTGRP/GGPGGPGMG KPRCF/RGEVFGIVIRAGSPGPG TGPGRG\GEAAGAKAED*E\WM PV\TKLGRLVKDMKIKSLEEIYL FSLAIKKSEIMSFLLGGLLSRDE VFE*LCPVQEQ\TRAGQ/RATRF KAFVALTTLGTTNG\HVGLGC* VPPREVATGNSWGPFILGQALH SSPVRRGLLGGTKNSAKPPTLV PWQR*PRAACGLCAGYAFIPLQ PRGQLASVSA\PVP\KKLLMMA GTDG\CNNSQRRGCTATVG\NL AKATTFDAISKTYSYLTPDLWK ETVFTKSPYQEFTDHLVKTHTR VSVQRTQAPAGATT
12982	43350	A	13062	92	265	SFELFADKVPKTA/WLDGKHVV FGKVKEGMNIVEAMERFGSRN GKTSKKITIADCGQLE

SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			sequence		
43351	Α	13063	2	797	NRVLLAMVNPTVFFDIAVDGEP
					LGRVSFEVRGLDTKK*LLI*SIK
					LC*QIG\LFADKVPKTAENFR/A
					L*SIEEKGFGL*GVPCFHRLFPR
					VLCVRGGDFHTAINGT\GGKSI\
	ļ				YGEK\FEDENFI\LKHTGPGILSH
					GKMLDPNTNGSQFFICTAKT\E
	l				WLDGKHV\VFGKVKEGMNIVE
	1				AMEA\FGSRDCKTSKKITIA\DF
]				GQLRIKFDFVFLSLTTKIISFCAL
	i				LSGEHPLHPIWLGVILESMWLS
					LAVPFWVPCFPCSLPCLAGLQS
43352	A	13064	1	720	
43353	В	13065	98	2074	
	Α		596		
43355	Α	13067	66		
43356	Α		1		
43357	Α	13069	2	325	WCPSQAAAAGARATRDMPEGS
	l				MTCLCRALWWLFSWLKVHR*R
	ŀ				QGQQGRPAAPSAGPAKPTPTRN
					SSWLLLAAPVPTRTSPSTPPCKL
					REPAPALASPERGSHSAAVG
43358	A	13070	38	1066	
43359	В	13071	1	324	
43360	Α	13072	2	680	
43361	A	13073	2	165	
43362	Α	13074	1	1110	MASLLKSARPQTHWKEETPETS
					KRLKEQTPDIPSLRAVTLIAKVH
]				SFIPEVCETKNPLEGIHSGHILAP
					NVGPSPSRRQARVQVFENVSVR
					ATKSDLPQSSLWSRRKTTVSAA
					ASKKTSKEISKGPQKPPGY/PVT
					SPSSCRGRGIWPNP/EYMSTSPS
					LI*/SQIKVDLGKISDDPDRYIDV
					LQGLGQTLDLSWRDVMLLLDQ
					TLAFNEKNAALAAAQEFGDTW
					YLSEVNDRMTAEERDKFPTED
					GEPIKCDCQQIIVQTYAAQDDIL
					EVPLANPDLNLYTDGSSFVENG
					IRRAGYAIVSDVTILERPNLFRA
					IQQVVKACEVCQRKNPLVHPP
					NYLYKTLNLTHSLLNQSNPSLA
					NDCWLCSSLSVSAEPYN
	43351 43351 43352 43353 43354 43355 43356 43357 43358 43359 43360 43361	43351 A 43351 A 43352 A 43353 B 43354 A 43355 A 43356 A 43357 A 43358 A 43359 B 43360 A 43361 A	of peptide sequence hod in USSN 09/540,217 43351 A 13063 43352 A 13064 43353 B 13065 43354 A 13066 43355 A 13067 43356 A 13068 43357 A 13069 43358 A 13070 43359 B 13071 43360 A 13072 43361 A 13073	Sequence 09/540,217 codon for peptide sequence 43351 A 13063 2	of peptide sequence hod in USSN 09/540,217 location of first codon for peptide sequence codon for peptide sequence 43351 A 13063 2 797 43352 A 13064 1 720 43353 B 13065 98 2074 43354 A 13066 596 841 43355 A 13067 66 326 43356 A 13068 1 441 43357 A 13069 2 325 43358 A 13070 38 1066 43359 B 13071 1 324 43360 A 13072 2 680 43361 A 13073 2 165

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12995	43363	Α	13075	450	1776	RGGRACLRP*AA/SPGPSTP*WP
						PPRGVQAAAILSGPPRPAPPKG
						AASQRLCGGFVEARQAWVGRR
						GGRRGPVGETQPPPSVIQPPSRG
						AG/PRDQQPAMPEPPTPSVGSC
		ŀ				AARASQISAAPCSTAPSPIDHPR
						AEECRRTVWDWQAAPPAAQV
						RDPLGEASWAPESGGDVENLY
						VLLRDCKYTSQHPVSSSGSVNA
						PIDTLYLATLLRDCKYTNRHSV
ł]				SSSRFVNTPISTVSSSEFANAPID
						TLHLATLKEETPNTSEHQKEQT
				1		PDMPPLRTVTVTARVRGFILDV
						SETKNPPIPDTFWRPRWDLRQS
						PSNQTLAFNEKNAALAAAQEF
						GDTWYLSQVNDRMTAEERDKF
		ŀ				PTEFIPSSGFLVLLTSRMKPQTF
						AASVTALKDGVSRVCSFRCVQS
						FFLPSADWCIYKPLARYRALIG
						AFLQSAHWCVYKPLARHSVLI
		<u> </u>				GAFTNL
12996	43364	В	13076	1	1999	
12997	43365	Α	13077	3	469	
12998	43366	A	13078	224	410	
12999	43367	A	13079	478 55	675	
13000	43368	C	13080	ļ	237	
13001	43369	A	13081	109	239 88	
13002	43370	A A	13082	2	178	
13003	43371	C	13083	278	460	
13004	43373	C	13084	50	217	
13005	43374	A	13086	3	251	
13007	43375	A	13087	1	759	
13008	43376	A	13088	2	604	
13009	43377	A	13089	1	558	
13010	43378	A	13090	3	1055	
13011	43379	A	13091	1	466	MDKFLDTYTLPRLNQEEVESLN
						RPITGAEIVAIINSLPTKKSPGPD
						GFTAEFYQRLISNFSKVSGYRIN
						VQES\QAFLYTINRQTESQIMSA
			-			LPLTIASKRIKYLGIQLTRDVKD
						LFKENYKPLLNEIKEDTNKWK
]			NIPCSWVGRINIVKMAILPK
13012	43380	Α	13092	2	848	
13013	43381	Α	13093	458	625	
13014	43382	Α	13094	482	745	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13015	43383	A	13095	1	742	MNIDAKILNKILANRIQQHIKKL
						IHHDQVGFIPGMQGWFNIRKSI
						NVIWHINRTKDKNHMIISIDAE
		1				KVFDKIQQPFMLKTLNKLGIDG
						TYLKILRAIYDKPTANIILNGQK
						LEAFPLKTGTRQGCPLSPLLFNI
						VLEVLARAIRQEKEIKGIQLGKE
						EVKPSLFADDVIVYLENPIISVQ
						NLLKLISNFSK\IKYLGIQLTRNV KYLFKENYKSLLNEIKEDTNK
						WKNIPCSWIGRINIVKMAILPK
13016	43384	A	13096	1	1458	
13017	43385	A	13097	2	1624	
13018	43386	A	13098	1	1095	
13019	43387	A	13099 13100	1	2913 1887	
13020 13021	43388	A	13100	1	972	
13021	43390	A	13101	1	873	
13022	43391	A	13102	1	1365	
13023	43392	A	13104	1	867	
13025	43393	A	13105	1	591	
13026	43394	A	13106	1	990	
13027	43395	Α	13107	1	2109	
13028	43396	A	13108	1	843	
13029	43397	Α	13109	1	1203	
13030	43398	В	13110	1	1221	
13031	43399	Α	13111	1	1092	
13032	43400	Α	13112	1	931	
13033	43401	Α	13113	1	1293	
13034	43402	A	13114	1	544	
13035	43403	A	13115	3	718	A PARTICULAR PARTICULAR PROPERTY AND A PARTICULAR PARTI
13036	43404	A	13116	693	1397	ARAEVKLSLFADDMIVYLENPII
						*ARAEVKLSLFADDMIVYLENP IISAQNLLKLISKFSKVSRYKINV
						QKSQAFLYTNNRQTESQIMSEL
			}			PFTIATKRIKYLGIQLTRDVKDL
		1				FKENYKPLLNEIKEDTNKWKNI
						PCSWIGRINIVKMAILPKVIYRF
		1				SAIPIKLPMTFFTELEKKNWLAI
						CRKLKLDFFFIPYTKINSRWIKD
						LNVRPKTMKTLEESLGNTIQDI
		İ				GIGKDFMTKTPKAMATKAKRA
						SAQQKKLPSE
13037	43405	A	13117	1	2814	
13038	43406	A	13118	1	1302	
13039 13040	43407 43408	A	13119 13120	3	1542 1327	
13040	43409	A	13121	1	1023	
13041	43410	A	13121	1	1448	
13042	43411	A	13123	1	1038	
13044	43412	A	13124	3	1711	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		07/340,217	sequence	or peptide sequence	defending to possible nucleotide insertion)
	<u> </u>				 	
13045	43413	В	13125	<u>l</u>	1756	
13046	43414	A	13126	1	923	
13047	43415	В	13127	1	1515	
13048	43416	В	13128	<u>l</u>	1560	
13049	43417	Α	13129	1	4032	
13050	43418	Α	13130	1	2684	MVKGSIQQEELTILNIYAPNTG
						APRFIKQVLSDLQRDLDSHTLI
						MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQADLIDINRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQSRSTPWKLN
						NLLLNDYWVHNEMKAEIKMFF
						ETNKNKDTTYQNLWDAFKAV
						CRGKFIALNAYKRKQERSKIDT
						LTSQLKELEKQEQTHSKASRRQ
						EITKIRAELKEIETQKTLQKINES
1						RSWFFERINKIDRPLARLIKKKR
						EKNQIDTIKNDKGDITTDPTEIQ
						TTIREYYKHLYANKLENLEEM
						DTFLDTYTLPRLNQEEVESLNR
1						PITGSEIVAIINSLPTKKSPGPDG
						FTAEFYQRAIYDKPTANIILNGQ
						KLEAFPLKTGTRQGCPLSPLLF
		İ				NIVLEVLARAIRQEKEIKGIQLG
İ						KEEVKLSLFADDMIVYLENPIV
ŀ						SAQNLLKLISNFSKVSGYKINV
						QKSQAFLYTKNRQTESQLMSEL
İ						PFTIASKRIKYLGIQLTRDVKDL
						FKENYKPLLNEIKEDTNKWKNI
						PCSWVGRINIVKMAIL\PKVIYR
						FNAIPIKLPMTFFTELEKTTLKFI
						WNQKRARIAKSILSQKNKAGGI
	}	Ī				TLPDFKLYYKATVTKTAWYW
						YQNRDIDQWNRTEPSEIIPHIYN
						HLIFDKPEKNKQWGKDSLFNK
						WCWENWLAICRRLKLDPFLTP
12051	12410	<u> </u>	12121	1	1140	W C W EN W LAICKRENDFFE IP
13051	43419	A	13131	127	1149 329	
13052	43420	A	13132	127		
13053	43421	A	13133	1	1132	
13054	43422	A	13134	1	1020	
13055	43423	A	13135	803	2009	
13056	43424	Α	13136	1	1944	
13057	43425	Α	13137	1	1282	
13058	43426	Α	13138	1	2423	
13059	43427	Α	13139	1	2694	
13060	43428	Α	13140	1	1195	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13061	43429	Α	13141	6082	6984	KLAQDRDALSPPLLFNIVLEVL
					1	ARAIRQEKEIKGIQLGKEEVKLS
						LFADDMIVYLENPIV\SAPKSPL
						RLISNVS\KV*GYKINVQKS\QA
						FLYTNNTDKQESQIMSELPFTT
						ASKRIKYLGIQLTRDVKDLFKE
						NYKPLLKE\IKEDTNKWKNIPCS
						WVGRINIMKMAILPKVN\YRFI
		İ				NSMPSIPIKL\PMTFFTELEKNYF
						KVHMEPKKSPHCQVNPKTKEQ
İ						SWRHQRYLTSNYILQGYSNQK
						STISMVLVPKQHGTGTKTEIIDQ
1						WNRTEPSEITPHIYNYLIFDKPL
		1				LEKNKQWGKGFPI
13062	43430	Α	13142	193	335	MVEEKLTLPKDEK*KSKQRSIY
						RKTLQELKNKNEKRNRKNKW
						QVKTH
13063	43431	Α	13143	131	427	
13064	43432	Α	13144	1	360	
13065	43433	Α	13145	2	376	
13066	43434	Α	13146	1	360	
13067	43435	A	13147	2	376	
13068	43436	Α	13148	3	469	
13069	43437	Α	13149	224	410	
13070	43438	Α	13150	478	675	
13071	43439	Α	13151	32	161	
13072	43440	Α	13152	1	1671	
13073	43441	A	13153	1	1656	
13074	43442	A	13154	62	1295	
13075	43443	В	13155	149	320	
13076	43444	A	13156	12	89	
13077	43445	A	13157	28	417	
13078	43446	A	13158	190	440 553	EIVDECA SESIDIH DWSDIIDVC
13079	43447	Α	13159	190	333	EIVREGASFSIRHLRWSRHRKC
						RFSVKTLTGEDPSTLEV\EPS\DT
		1				IE\NVKAK\IQDKEGI\PPDQQRL/
						VSFAGKQL\EDGRDSLSGY\NIQ
						KGSLLLHPCV*DFRGGAKKKEE
12000	42440	<u> </u>	12160	2	415	ESLYTTSPRKD
13080	43448	A	13160	3	415	PVK/VGA*GGQVINGVLAQV*L
						TVGPVGPRTHPVVIFPVPECRIG
						RDILSSWQNPHTGSLTGRVRAF
						MVGKAKWKP*ELPLPRKTVNQ
			1			KQYRIPGGIAEIS/A/TIKNLRRG
					1	VVIPTTSRFNSPIWPVQKTDGS
		<u>l</u>	L	<u> </u>	<u> </u>	W*TAADY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
	÷	<u>ļ</u>			ļ	
13081	43449	Α	13161	5	405	HCCGIPHSIASEQDTYFMAKEV
					•	WQWAHAHGIHWSYHVPHHLE
						AAGLIEQWNGLLMSQLQHQLG
1			i			DNTLQGWGKVLQKVVYALNQ
						RSIYGTVSPIARIHGSRNQGVEV
	·		i			EVALLTVTPNDPL/GKY*LPVPV
						TLHSDR
13082	43450	A	13162	138	387	
13083	43451	C	13163	184	297	
13084	43452	Α	13164	1	430	
13085	43453	Α	13165	1	867	
13086	43454	Α	13166	35	601	CLSRRASPVYLASMSGRGKTG
						G\KARA\KAMSRSSRAGRPSSQ
]	VGR\VHRLLRKGHYAERV\GRR
						QPCCYLGG*CMEYLTS*ESWS
						1 ~
						MAG\NAARNN\KKTRIIP\RHLQ\
	1					LA\IRNDEELNKLAWAALTIAQ
						GSR/VLPNIQARCCGPR*TSATV
						G\PKAPSGGEERATQASARSTK
						RARAAAGRPSVPMPPQRPF
13087	43455	С	13167	11	313	
13088	43456	В	13168	1	552	
13089	43457	Α	13169	1	621	
13090	43458	A	13170	55	296	
13091	43459	Α	13171	959	1182	WVYLF*PSCKGVIYP*AAGFESI
						FWWVTINKNIDWINYIYYNQQ
		}				QFMNYTRDAVKGIAEQLGTNC
		1				QMAWENRIALDMILAERGGVC
						IMIKTECCAFIPNNTAPNGSITK
		1				ALQGLTALSNELASSSGVNDPF
						TGWLEKWFGKWKGITASILTSL
						TAVMGVLILVGCCVIPCICGLV
	İ					QRHRGPPLVVIETKPLGLERLA
						1 -
		1				GLPVGHALKLGSGIQATPQNGE
12002	43460	<u> </u>	13172	1	540	NADRREAFPAAS
13092 13093	43461	A	13172	2	142	
13093	43462	A	13174	42	384	
13095	43463	A	13175	102	329	
13096	43464	A	13176	1	1257	
13097	43465	A	13177	1	378	
13098	43466	A	13178	1	1143	
13099	43467	A	13179	1	481	SDSELNRPRVTMKFNPFVTSDR
		1				SKNRKRHFNAPSHVRRKIMSSP
						LSKELRQKYNVRSMPIRKTDEV
						\QVVRGHYKGQQIGQGSPRCTR
						KKYVIYIER\VQREK\ANGTNCS
						~
						TWGISPKARWLFTRLKLDKDR
						KKILERKAKSRQVGKEKGKYK
1		<u>L</u>				EELIEKMQE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13100	43468	A	13180	39	556	AAAGYFAEDCEASCVCVKHPP
						SVKKARCFLSELIKKPSGGSVTL
						SESTAIISHGTTGLVTWDATLYL
ŀ						AEWAIENPAAFTNRTVLELGSG
						AGLTG/VAICKMYRPRAFIFSDC
						HSRVLEQLRGNVLLNGLSLEAD
						ITANLDSPRVTVAQLDWDVAT
						VHQLSAFQPDVVIAADVLY
13101	43469	Α	13181	46	231	
13102	43470	Α	13182	1	1503	
13103	43471	Α	13183	3	340	FFFANTFYCVFNVLVNAPLRFL
			,			SLPSTQSLEAKLRDSSDSELRRD
Ì						ILQKVRIPESLGLMTLPPESLRK
						T*EKNRRCACTL*K\TPVQRTKP
						LSQHCRWVWAPPTAPALSFSKT
13104	43472	Α	13184	2	594	
13105	43473	Α	13185	1	1470	
13106	43474	В	13186	79	1355	
13107	43475	Α	13187	248	540	
13108	43476	Α	13188	386	1388	
13109	43477	Α	13189	386	1334	
13110	43478	Α	13190	3873	4070	
13111	43479	Α	13191	146	510	PPCAVVCLLWIAAVYCLSTTFC
						APLCGQNTWLPKPCRHVNRAA
				1		SNASVSLSTVTLFSRLS*PQRKG
		İ				SELLDSSGPLPASPSLPLCSGVS
						PRAGLGSAWPKIPFLGIREAKN
						PRSENTRLTT
13112	43480	Α	13192	21	95	KCSSNGAPDAVHE*DLPWTPGP
						AC
13113	43481	В	13193	1	477	
13114	43482	Α	13194	228	935	PLSSSMAAELEFAQIIIIVVVVTV
						MVVVIVCLLNHYKVSTRSFINR
						PNQSRRREDGLPQIMHAPRSRD
						RFTAPSFIQRDRFSRFQPTYPYV
						QHEIDLPPTISLSDG\EEPLLYQV
						PCTLQLRDPEQQMELNRESVRA
						PPNRTIFDS\DLIDIAMYSGGPCP
						PSSNSGISASTCSSNGRMEGPPP
	1					TYSEVMGHHPGASFLHHQRSN
						AHRGSRLQFQQNNAESTIVPIK
L						GKDRKPGNLV

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	1	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13115	43483	A	13195	526	1629	FYSFDSARGEPARPWRVRGLEG RVELGQQLGLLTLGPAFPGGLG GTQRVGLPGPAGPGAGEEGQA GPAEAGQVPVGGLADEAVEHG VGDAVEAGEQEGEVVGVENAL GEVAAGLPDAAHQQQHVVGQ EAGQEDDHGAEHQPLDLVLAA LLGAVAPAHGAQDALVGRQQ QADGEEEA/LPGSGSS*WHAAM PGWGSAQSTRGAGSRPR*PPRA SGSRCWSAAPWPR*PR*RRPP SGSGTGGSGRDGPQPGSGPR/T MQQRKPTLM*MFSYRR\IRTPG RTGRRETSCRAAARTAARRAA SRCRRGRPPSGCTGTRPARSGP* SAGS\EVEGQAVGREAHHQHG NIDHRGQRLVDGMVDGTAHRG VVCSDVPH
13116	43484	В	13196	2699	7361	VVCSDVIII
13117	43485	A	13197	67	1523	
13118	43486	A	13198	1	287	
13119	43487	A	13199	2	563	
13119	43488	A	13200	96	401	
13121	43489	В	13201	1	480	
13121	43490	A	13202	1	924	
13122	43491	B	13202	13	1512	
13123	43491	A	13203	1	402	
13125	43493	A	13205	609	803	RDPRHNPSSA\FQAGGIALLMLI TRGEDLTGGVIHQYPGPGMNL WMQGTAGNQRRAAENAAQQQ
13126	43494	Α	13206	3	526	VTETALTLPYLVAKHGYFLPQD LAKRTETMNWLFWLQGAAPFL GGGFGHFYHYAPVKIEYAINRF TMEAKRLLDVLDKQLAQHKFV AGDEYTIADMAIWPWFGNVVL GGVYDAAEFLDAGSYKHVQR WAKEVGERPAVKRGRIVNPTN GPLNEQLHERHDPSDFQTNTE\ NNRQG
13127	43495	Α	13207	1	250	
13128	43496	Α	13208	1	489	
13129	43497	Α	13209	209	601	
13130	43498	A	13210	1	96	
13131	43499	Α	13211	1	1110	
13132	43500	Α	13212	1	595	
13133	43501	A	13213	1	240	
13134	43502	A	13214	1	675	
13135	43503	Ā	13215	1766	1957	
13136	43504	A	13216	1	426	
13137	43505	В	13217	94	3238	
13138	43506	A	13218	1	2370	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13139	43507	A	13219	1	2796	
13140	43508	В	13220	43	566	
13141	43509	Α	13221	1	903	
13142	43510	Α	13222	3	110	
13143	43511	Α	13223	2	755	
13144	43512	Α	13224	1	1338	
13145	43513	Α	13225	1	807	
13146	43514	A	13226	1	918	MARADTVSVPMFMGLAAKPC WRDTEPNTGYRGPHVRNIQLT
		1				HDPRLDYRSI/LIDINDIGQTFHE
						RLHPDAC\LSNAILVHNKKGGP
			ŀ			LADGIVITPSHNPPEDGGIKYNP
						PNGGPA\VPTSLKWWKTGPTHC
			ŀ			WPMA*KA*SVSP\RRSDGIRSKN
ĺ						DVIDTSDDDM**QCLSRRSN\G
						HPGAKVANLVPKTTFIDGVDQT
						SFFLGTNGQSNRKAEHYFLNGK
						LAAVRMDEFKYHVLIQQPYAY
						TQSGYQGGFTGTVMQTAGSSV
						FNLYTDPQESDSIGVRHIPMGV
13147	43515	A	13227	1475	2904	PLQTEMHAYMEILKKYPPRAQI FWRAAAPIPDCWRSGAVKKQH
1314/	43313	A	13221	1473	2904	VEVTQLDWTPPGRQYAGPIPCS
					1	RRGYCPLPRSGSVRWADL/FPP
						A/LD/LAGHPG/AKVANLVPKTT
						FIDGVDQTSFFLGTNGQSNRKA
						EHYFLNGKLAAVRMDEFKYHV
						LIQQPYAYTQSGYQGGFTGTV
	!					MQTAGSSVFNLYTDPQESDSIG
						VRHIPMGVPLQTEMHAPGLPLA
	ļ					SSMKRWIRRATGFIFGKEQFED
		1				VVPVLGSKVNGVQFNAELVAD
						SLGISQIRCRCAIFLTVVFFPVLH
						KQAFDLISLLLQQPGRNGGIDT
	L					AGHADDYFFCGFRHWITHDIEL
	i i					QGGARQKDAADHYLAGDIESL
						PLATATFDLAWSNLAVQWCGN
						LSTALRELYRVVRPKGVVAFTT
		1				LVQGSLPELHQAWQAVDERPH
		1				ANRFLPPDEIEQSLNGVHYQHH
		İ				IQPITLWFDDALSAMRSLKGIG
						ATHLHEGRDPRILTRSQLQRLQ
L		1				LAWPQQQGRYPLTYHLFLGVI
13148	43516	Α	13228	215	525	LAWARRCRRLLAKATSLIRAPS
						SEMRASTPCRGQRRPSGYSWY
						GRSPRAVLEGLRVGAHLSRAN
		1				CPWPSR/SQVVELIPDGFPVFLD
						FTKIPGRDTQNEHIVLLH
13149	43517	Α	13229	1	942	
13150	43518	В	13230	1	1201	
13151	43519	Α	13231	685	921_	

13152 13153 13154 13155 13156 13157	SEQ ID NO: of peptide sequence	A A A A A	SEQ ID NO: in USSN 09/540,217 13232 13233 13234 13235 13236 13237	Nucleotide location of first codon for peptide sequence 2 2 2 5 2 1 572		Amino acid sequence (X=Unknown, *=Stop codon,/=possible nucleotide deletion, \=possible nucleotide insertion) RCRPDKAFTPHPAIGARCLMRR LRVLSCLQPLPNVGWIRRLRRIR
						QFSAPDEGAKCRVAILQ/DFDA QYGRFLEVTSGAQQRFEQTD*K HAAALPFFFPSSAYSLAAISVVF GFR
13158	43526	C	13238	5	127	
13159	43527	Α	13239	1	444	
13160	43528	Α	13240	17	261	
13161	43529	Α	13241	173	397	
13162	43530	A	13242	169	421	LKPKNLDEKLLPASSSSCRIWA TSPVHHLWQVLKKILFGWNPT KSPRCLS/TRQAMFSLSKKASLP WTIAARSADQRCRTRIAEL
13163	43531	A	13243	17	744	RPHQEQRIAGCGVNALSSLRSA TAVGMIRRASVASG/DRAPIAG CGVNALSGLHRHFIEVTCTHFL LVRYEGIAIFRGGKFRLLHFLN VVLHTLALGIGVCQVKHVEPH AVDTCQGDELELVAHIRQLLLE AGNSFVVEVYLPVERRAVISQ QFARIFRVDSLCKATRQFQIRRG RSHQTGQHMGIRRQVDAAHAD DRRTSGRVPPLIQVCPAHSSEA VVVVDVDVDVDVDVDVDVIGAV
13164	43532	Α	13244	954	1104	
13165	43533	В	13245	54	1169	
13166	43534	A	13246	35	3066	VGHSTRPAASPGPAAGIGRSPA RLVQQPQRSLAKPLNRPAQQR GLLPQRRHPKRTRKRRKQAQN PQKRLPHRQPVRRRHRHHRRLL QKMRRPDKRQQRRAAPRRHPR RRQRLLAVRRRQLRAKVRRNP RQRAPRQQLNGQRILHPPWRLR MQVRRKRG*YSSAVRPTVRLK RWRQRQRQ*NQPMTMQRNVC RKTRTALIYP/YKGCFLNNINAV SKTDFADKRGMRYVRVNAPAG ATSGKYYPVVVMRSAGSFHI
13167	43535	A	13247	1	2919	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13168	43536	A	13248	1	3211	MEFTSLTPDDLRVLELYRTPFS
						DGYVFHSMQFHIFDLLKSGQN
				}		VVLSAPTSMGKSAIVDSLLGMG
						TLKRLVLVVPTVALADETRRRL
						QERFGDRYQIIHHSSQVCHSDQ
		1				AVYVLTQERVNERDDIVDIDLF
						VIDEFYKLAFRQLKSGDIDHQD
]				ERVIELNIALSKLLKVSRQFYLT
						GPFVNSIRGLEKLGYPHTFVST
						DFNTVALDVKTSASKRMTTKP
					•	SSKRWGKSRTCVDADHIKNGIG
						LHFGALPRALQQYT
13169	43537	Α	13249	486	3882	LRLRGEALPVASGCLANATDTP
						AGTTLWADTERHAVQ**TCEA
						DTLGSCGYAAKPCLSPVAVWQ
						MLLTRLLEQHYGLTLNDTPFSD
						EPVLTAPIDAAALRGIEFGKFQL
Ì			,			GNMVVGDIIECGSDVTDYAVG
						DSVCGYGPLSETVIINAVNNYK
						LRKMPQGSSWKNAVCYDPAQF
						AMSGVRDANGARRGFWWGGL
		l				GAIRVQLAFKWLNAAGRLPGG
						GLGGPIPYSAHRCDIARRHGAD
						FCLNPIGTDVGKEIKTLTG
13170	43538	В	13250	1	1116	
13171	43539	Α	13251	729	985	SAPPQLAPSLPDRATDPDTCWP
						PG*RP\RYLPSRQIDCFNTRFRLL
						QRLTASQCAEAVDVAFFRLTV
	<u> </u>	1				QQTPHFRCAQLRQRAFRID
13172	43540	Α	13252	395	605	
13173	43541	Α	13253	1	1140	MIYVFVTVKHRNDHFTKNTLA
					:	SYRGRTLLGTLFKGVYHLYKD
						ETYLYQSGKGHTIQEVRIVKGL
						NNPDLDAAVGEDLAQQLRDEL
				}		ELVKGASNEFDKELFLAGEITP
						VFFGTALGNFGVDHMLDGLVE
						WAPAPMPRQTDTRTVEASEDK
						FTGFVFKIQANMDPKHRDRVA
						FMRVVSGKYEKGMKLRQVRT
						AKDVVISDALTFMAGDRSH/AY
						PGDILGLH\NHGTIQIGDTFTQG
						EMMKFTGIPNFAPELFRRIRLK
			B	1		DPLKQKQLLKGLVQLSEEGAV
						QVFRPISNNDLIVGAVGVLQFD
						VVVARLKSEYNVEAVYESVNV
						ATARWVECADAKKFEEFKRKN
						ESQLALDGGDNLAYIATSMVN
12174	12512	_	12254	1	1405	LRLAQERYPDVQFHQTREH
13174	43542	A	13254	1	1495	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13175	43543	A	13255	372	1365	IRYFENRGLFAHCATRTGQFPP
					:	HCILASYKCGEQPDSSSNHPRN
						LFVWRSNLLGSSG\KGHEYMQ
						KYLLGTESGIQGEELGASDGIKP
						EEVEWQTAAIEGKLDLLVTLDF
						RMSSTCLFSDIVLPTATWYEKD
						DMNTSDMHPFIHPLSAAVDPA
						WESRSDWEIYKGIAKAFSQVCV
						GHLGKETDVVLQPLLHDSPAEL
						SQPCEVLDWRKGECDLIPGKTA
						PNIVAVERDYPATYERFTSLGP
						LMDKLGNGMDLTDYANMALSI
						PSANTDIWNLEQDTVGTRLTNS
						RHGLADNGGAWVSYFGGNFN
						GDNGTINYDQDVNGIMVGVDT KIDGNDG
13176	43544	Α	13256	3	1444	RIDONDO
13177	43545	A	13257	61	1058	LPYLIALLARAWFNGLLTSRTR
13177	143343	^	13237	01	1036	LYIKGNGIVDLPKLQEMVADVS
						HHFPLRLPAPTPKALYSPCEIRH
						LAINVNLEYDPTAAFRNQVVHF
						DFRKLDVFSFGENQNCLVGNV
						DLL\YRNSWNEVRTLHFNGEQS
						MIEALKTILGKMHQDAAPPDSV
						EVFCYSQHLRGLIRTRVQQLVS
		ŀ				ECIELRLSSTRQETGRFKALRVS
						GQTWGLFFERLNVSVQKLENAI
						EFYGAISHNKLHGLSVQVETNH
						VKLPAVVDGFASEGIIQFFFEET
						QDENGFNIYILDESNRVEVYHH
						CEGSKEELVRDVSRFYSSSHDR
						FTYGSSFINFNLQDPYLRARVSI
13178	43546	Α	13258	1	1133	
13179	43547	A	13259	2	240	ISSARHFGALACTLELGKALTF
						GQNDLRQFAVTASAIAALL/SR
						HSPSFEMHMASDTLNFMPFEK
						GTLLAQDGEERFTVTH
13180	43548	A	13260	1	358	LPFLPSGLRGLAAENKIRRPHSA
						GSFLGPPCGLRGKLV/ISAAALM
						STTIPISRVQGLLQFLNSKSPISR
						AERSWSTFGASAAFLTTLASLL
						FLFPPLAISSLANSFAKVFPPTPC
						AGSP
13181	43549	A	13261	1	1635	
13182	43550	Α	13262	1	1363	
13183	43551	Α	13263	5	249	
13184	43552	Α	13264	15	280	
13185	43553	A	13265	6	363	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	·	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13186	43554	A	13266	78	346	TRYAMLTRHDNHLSKFYLARG
13100	13334	l' `	13200	, ,		RYRADFHAPETNTGERCYRYK
						PLPYWLE*RRGRNRYPALQ*TM
						FLPADGPARCLSRQAAILKESV
						LPK
13187	43555	Α	13267	568	831	ELQRHVLVVFAGFWIVKNSRN
						LFLVCRAEHKRSVVKGLLRQQ
			1			GQGFWFHFQDGFAVEVGNAN
		1.		_		VIGSE*IVFGIVFPHRERCLVDK
13188	43556	A	13268	3	419	
13189	43557	A	13269	1	561	
13190	43558	Α	13270	1	741	
13191	43559	A	13271	2	1040	
13192	43560	В	13272	1	1974	EVELDOVI A CERTAIN
13193	43561	Α	13273	2111	2340	EVFIRDKLMERRNRRTGRTEKA
	1					/RDRTVRTWIGEAVAAAAADG
						VTFSVPVTPHTFRHSYAMHML
12121	10.7.60	<u> </u>	10071		0.10	YAGIPVKVLQSQCI
13194	43562	Α	13274	470	813	KTTSFPAKLFNACRISLLASAEV
						NPNISAYLATWISQIPSCSCRVP
						SDFLIWSTNRYSIRF*SRERLVG
						EAVA/AAAADGVTFSVPVTPHT
						FRHSYAMHMLYAGIPLKVLQS
10105	42.5.62		12075	110	1100	LMGH
13195 13196	43563 43564	C A	13275 13276	1	1180 1290	
13196	43565	A	13277	1	1036	MKLMETLNQCINAGHEMTKAI
13197	43303	^	13211	1	1030	AIAQFNDDSPEARKITRRWRIG
						EAADLVGVSSQAIRDAEKAGR
						LPHPDMENSRDRFEQRGFEPDV
						RILLTKYSNSNGSQSPWMEEQI
						RDAWGSMVLKNVVRETDEVG
						KDPRPLPYLGHDEPYTFDINLS
						VNLKSMVVGGTDMGQEPLKQ
						1
						MGAWGPLSLKAMFGFFIRGYG
						FFTPFGRTLFPLGFGSTPPFTFPL
						LTKTIGVFILDKLMERRNRRTG
	ŀ					RTEKARIWEVTDRTVRTWIGEA
						GAAAAADGVTFSVSVTPHTFR
						HSYAMHMLYAGIPLKDLQSLM
						GHKSISSTEGYTKDFSLDVG\AR
10100	10.055	ļ.,	10000		C 70	HRVQF\AMP\ESDAV\AMLKQLF
13198	43566	Α	13278	366	672	RNGTHPRYIEAVPWGALPAD/G
						VTFSVPV/TPHTFRHSYAMHML
				1		YAGIPLKVLQSRLALRELQQAV
						HAGLPQQAKILFDGGSEIGKIPH IVLYKPVQCSLWAFVH
13199	43567	A	13279	2602	2976	IVETRI VQCSEWAI VII
13200	43568	A	13280	982	1347	
13201	43569	A	13281	1	780	
13202	43570	Α	13282	568	891	
	1.22,0	1	1	<u></u>	<u> </u>	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13203	43571	A	13283		496	MKEKPSSPRQYSYSFLSSMILLG GGESWNLLRADQRLIFAKSWP RASRYQQGHQDLFILRSDLPSQ VFIRDKLMERRNRRTGRTEKAR IWEVTDRTVRTWIGEAVAAAA ADGVTFSVPVTPHTFRHSYAM HMLYAGIPLKVLQSLMGHKSIS STEVYTK\VFALDVAAPHPESTP VHLPSRENKIPPPIMRIPCGLPTS PIEQKQGKISATGTQWRRLKQE TRLSSVSARLMSYVGVMC AMFGFFIRGYGFFTPFGRTLFPL
		A				GFGSTPPFTFPLLTKTIGVFILDK LMERRNRRTGRTEKARIWEVT DRTVRTWIGEAGAAAAADGVT FSVSVTPHTFRHSYAMHMLYA GIPLKDLQSLMGHKSISSTEGYT KDFSLDVG\ARHRVQF\AMP\ES DAV\AMLKQLF
13205	43573	Α	13285	2	545	
13206	43574	В	13286	1	735	
13207	43575	A	13287	1	1599	
13208	43576	Α	13288	583	1005	
13209	43577	В	13289	31	1239	
13210	43578	Α	13290	1	1023	
13211	43579	Α	13291	1	933	
13212	43580	A	13292	1	1278	
13213	43581	Α	13293	1	705	
13214	43582	A	13294	2	893	
13215	43583	A	13295	88	429	
13216	43584	В	13296	1	1419	
13217	43585	Α	13297	1	2784	
13218	43586	A	13298	163	416	RNGTHPRYIEAVPWGALPAD/G VTFSVPV/TPHTFRHSYAMHML YAGIPLKVLQSRLALRELQQAV HAGLPQQAKILFDGGSEIGKI
13219	43587	A	13299	740	1431	CCQTLPVFHPPHPSHLGPRPCPP TH/HLPSEHRSLFLEAACHDSDS LEPLNLSSGSKTKSPSLPPKAKK PKGLEISA/RPAGALRHRHRLHR PQQPSPPLGIPHPSLL\TAQVFIR DKLMERRNRRTGRTEKARIWE VTDRTVRTWIGEAVAAAAADG VTFSVPVTPHTFRHSYAMHML YAGIPLKVLQSLMGHKSISSTE VYTKVFALDVAARHRVQFAMP ESDAVAMLKQLS
13220	43588	В	13300	1	858	
13221	43589	В	13301	1	1098	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13222	43590	Α	13302	1	1047	MNRQLSDSYTEDTKEPSDVTTS
		ŀ				ERTRSPPGSAKTTMIDTLKKLQ
						DVLKTEDSKNPTKSAADLLEQ
						YVKATGPIIEILQKATKTMEMI
						MIVEEKASDELQPDPELQQRYI
						HSVRGKPGLVKQRTQEIETRLR
			 			LAGLTVSSPLKRSHSLAKLGSL
						TFSTEDLSSRPPGPPPGPPPGLHP
			1			GPPPRRPPPKLPPPAPPGCSVHC
}						CHYPLHFVVSPLEGFYCSLMRI
		İ				QIVFVTAFVHHDSALLPLDASL
		ŀ	i i			APEALQNFQRYTGIQHVHRIGM
						AERMWCDRNRERHTVSSSGGN
			İ			RLPNPAPLPNPGVLSAPPNLIQR
ļ						PKADDTSAATIEKKATATISAK
ľ						PQITNPKAEIT*NPRVWQGSWI
						GEAVAAAAADGVTFSVPVTPH
						TFRHSYAMHMLYAGIPLKVLQ
						SLRSQ*SVRGKPGLVKQRTQEI
						ETRLRLAGLTVSSPLKRSHSLA
						KLGSLTFSTEDLSSRPPGPPPGP
						PPGLHPGPPPRRPPPKLPPPAPP
						GCSVHCCHYPLHFVVSPLEGFY
						CSLMRIQIVFVTAFVHHDSALL
						PLDASLAPEALQNFQRYTGIQH
						VHRIGMAERMWCDRNRERHT
						VSSSGGNRLPNPAPLPNPGVLS
						APPNLIQRPKADDTSAATIEKK
						ATATISAKPQITNPKAEITRFCPL
13223	43591	В	13303	1	1185	
13224	43592	A	13304	529	774	
13225	43593	В	13305	1	1089	
13226	43594	Α	13306	712	1024	GVESNLVVVVALCHRLIYLVW
1						GTRTVRTWIGEAVAAAAADGV
						TFSVAVTPHTFRHSYAMHMLY
						AGIPLKVLQS\RVVALDVAARH
12007	42505	_	12207	557	1040	RVQFAMPESDAVAMLKQLS
13227	43595	A	13307	556	1040	
13228	43596	A	13308	l o s	984	
13229	43597	A	13309	95	428	
13230	43598	С	13310	1	426	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13231	43599	Α	13311	116	1746	SSVTGRTEKARIWEVTDRTV/R
						RPWIGEAVAAAAADGVTFSVP
						VTPHTFRHSYAMHMLYAGIPL
						KVLQSLMGHKSISSTEVYTKVF
						ALDVAARHRNRFTQFRLSETKE
						ITNPYAMRLYESLCQYRKPDGS
						CIVSLKIDWIIERYQLPQSYQLY
						YFELAIPVGYFYPGSFSTASRIL
						LLHPRGLRAITIAVFGKQNTYIR
						LEPFKINVLEQITKHIEKLQCGG
						VVKQLSRRGNNQHISSTYDINR
						ADTQVRRAVNNYDIIVMSNSFN
						GQSEHQVWIARLTWVIGTINVV
						CAADVLIVPTPAELFDYTSALQ
						FFDMLRDLLKNVDLKGFEPDV
						RILLTKYSNSNGSQSPWMEEQI
		ŀ				RDAWGSMVLKNVVRETDEVG
					·	KARLTWGIGTINVVCAADVLIV
						PTPAELFDYTSALQFFDMLRDL
į						LKNVDLKGFEPDVRILLTKYSN
	1		•			SNGSQSPWMEEQIRDAWGSMV
						LKNVVRETDEVGKEPPSTNTFR
						HSYAMHMLYAGIPLKVLQSLM
					·	GHKSISSTEVYTKVFALDVAAR
						HRVQFAMPESDAVAMLKQLS
13232	43600	A	13312	1	1593	
13233	43601	Α	13313	1	2536	
13234	43602	Α	13314	887	1205	RPEKAR/IWGVTDRTVRTWIGR
		ŀ				AVAAAAADGVTFSVPVPPHTF
						RHSYAMHMLYAGIPLKVLQSL
						MGHKSISSTEVYTKVFALDVAA
						RHRVQFAMPESDAVAMLKQLS
13235	43603	Α	13315	79	351	
13236	43604	Α	13316	1	2320	
13237	43605	A	13317	187	798	
13238	43606	Α	13318	2089	2610	
13239	43607	Α	13319	1	1407	
13240	43608	В	13320	154	3682	
13241	43609	A	13321	1	1206	
13242	43610	A	13322	1	255	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13243	43611	A	13323	3058	3160	GSAA*LPPPLPAAFSGSCWLSQ RHPELHV*NEIYHLLH*KSFTSC *HNLSS*PRYPYLSSERVLW*CL SFISKI*SVTMP*SDLISLW*KLE PLTCRSTSHFRQKLAQ/RLP/AY QQGHQDLFILRSDLPSQVFIRDK LMERRNRRTGRTEKARIWEVT DRTVRTWIGEAVAAAAADGVT FSVPVTPHTFRHSYAMHMLYA GIPLKVLQSLMGHKSISSTEVYT KVFALDVAARHRVQFAMPESD AVAMLKHYPEINALALYLKCG TEWIC
13244	43612	Α	13324	1	1812	
13245	43613	В	13325	1	1776	
13246	43614	Α	13326	583	751	
13247	43615	В	13327	826	2097	
13248	43616	A	13328	1810	4163	
13249 13250	43617	A A	13329 13330	1085	2939 1690	SYFIMVKVGTSYVPINVSFSPKV
						GPGASRYQQGHQDLFILRSDLP SQVFIRDKLMERRNRRTGR\IYK ARIW\EV\TDRTVR\TWICRGGL PAAAADGVTF\SDPGPP\HTFRH SYAMHMLYAR*YR*KFLQSLM GHKSHPINGKATPKVFGPGMG LARA/HRGAGWQWPKSDAVG\ MPKQLSLRINALGLYMETGKPE GIMLFFGC
13251	43619	A	13331	3	327	
13252	43620	A	13332	3	337	
13253	43621	C	13333	167	1256	
13254 13255	43622 43623	A	13334	1	1078	
13256	43624	A	13336	3	976	
13257	43625	A	13337	1	1004	
13258	43626	A	13338	2	291	WRKIYEANGKKRKQRLQS*SL MKQTLNQQRSKETKKASA/SM RIKYLGIQLTRDVKDLFKENYK PLLNEIKEDTNKWKNIPCSWVG RINIVKMAIPPK
13259	43627	Α	13339	1	2022	
13260	43628	Α	13340	1	2262	
13261	43629	Α	13341	5	283	
13262	43630	A	13342	1	1245	
13263	43631	Α	13343	1	1203	
13264	43632	Α	13344	1	687	
13265	43633	В	13345	1	1371	
13266	43634	Α	13346	1	2241	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13267	43635	Α	13347	1	832	LKKCIKTPEIPSKHVRNWVPKV
						LEQRRQGLETYLQVASITLIPKA
						DEDTPKKKTTDKNSWTGTDAK
						VFNKILEIKFSSILKRLYWPVLE
						VLARAMRQEKEIKGIQLGKEEV
						KLSRFADDMIVYLENPIVSAQN
						LLKLISNFSKVSGYKINVQKSQ
						AFLYTNNRQ\TESQIRIKYLGIQ
						LTRDVKDLFKENYKPLLKEIKE
						DTNKWKNIPCSWVGRINIVKM
		ł				AILPKRYPFCQLCHQSAISTTKG
						VFTSILPLLMIEQLVSGRSCPSV
		ŀ				AWSSMECTTV
13268	43636	Α	13348	1	1314	
13269	43637	Α	13349	1	843	
13270	43638	В	13350	1	1011	
13271	43639	A	13351	1	173	
13272	43640	A	13352	1	323	
13273	43641	A	13353	3	2020	
13274	43642	A	13354	1	1473	
13275	43643	В	13355	1	636	
13276	43644	В	13356 13357	1 1,	456 619	
13277 13278	43645 43646	A	13358	1	1293	
13278	43647	В	13359	79	1041	
13280	43648	A	13360	3	718	
13281	43649	A	13361	1	612	
13282	43650	A	13362	1	992	
13283	43651	A	13363	1	1236	
13284	43652	A	13364	1	1071	
13285	43653	A	13365	359	1250	KLPMGVAVVKLLHLPKVIIVFW
						LASWMMSRWLHYPDWVTIIAQ
						QVLMNVVSVMQADCRMNLLEI
						FLRWLMRKIFHVRLLPAVSTPP
		:				NCLNQLLLFFLTPVNYLPGQVM
						HFKKPLQIKRNYLSSRHLTFMS
						RKKLGKVSG\YKIHVQKS\QAFL
						SANSGQTESQIMSELPFTMASK
						RIKYLGIQLTRDVKDLFKEKYK
						SLLNEIKEDTNKWKNNPCSWV
						GRINIMKMAILPKVIYRFNAIPI
						KLPMTFFTELEKTTFKFIWNQK
						RAHIAKSNLSQKNKAGGITLPD
			<u>L</u>			FKLYYKATVTKTT
13286	43654	Α	13366	1196	2132	
13287	43655	Α	13367	1	2271	
13288	43656	Α	13368	2	1105	
13289	43657	A	13369	1	2117	
13290	43658	A	13370	1	1447	
13291	43659	A	13371	1	1669	
13292	43660	Α	13372	<u> </u>	2823	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13293	43661	Α	13373	1089	2539	
13294	43662	Α	13374	1	1787	
13295	43663	Α	13375	1205	1207	
13296	43664	В	13376	1	2745	
13297	43665	Α	13377	1	2063	
13298	43666	Α	13378	1	4380	
13299	43667	A	13379		3347	MGDFNIPLSTLDRSTRQKVNKD TQELNSALHQADLIDSYRTLHP KSTEYTFFSAPHHTYSKIDHILG SKALLSKCKRTEIITNYLSDHSA IKLELRIKNLTQNCSTTWKLNN LLLNDYWVHNEMKAEIKMFFE TNENKDTTYHNLWDAFKAVCR GKFIALNAHKRKQERSKIDTLT SQLKELEKQEQTHSKASRRQEI TKIRAELKEIETQKTLQKINESR SWVFERINKIDRPLARLTKKKS EKNQIDAIKNDK
13300	43668	Á	13380	1	2367	
13301	43669	A	13381	1	1116	
13302	43670	A	13382	1	1095	
13303	43672	A	13383	670	3131	QTESSSTSKSLSTMIKWASSLG WQGWFNIRKSINVIQHIKGIQR QKPHDYLNRCRKSL*QNSTTLH AKNSQ*IVAQNLLKLISNFNKVS GYKINVQKSQAFLYTSNRQTES QIMSELPFTIASKRIKYLGIQLTR DVKDLFKENYKPLLKEIKEDTN KWKNIPCSWVGRINIVKMAILP KVIYRFNAIPIKLPMTFFTELEK TTLKFIWNQKRAHIAKSILSQK NKAGGITLPDFKLYYKATVTKT AWYWYQNMCYRSMEQNRALS YYVAYLQPSDL
13304	43672	Α	13384	441	3131	
13305	43673	В	13385	1	5688	
13306	43674	В	13386	916	2165	
13307	43675	A	13387	1	1529	MDPPAGAARRLLCPALL*LPAG *PLRPRLAGAAPAPGTALLHER AMPL*LRPGLHDVHGVPDL
13308	43676	Α	13388	1	714	
13309	43677	Α	13389	3	457	
13310	43678	A	13390	2	492	
13311	43679	Α	13391	3	1201	
13312	43680	Α	13392	1	309	
13313	43681	Α	13393	2	1073	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13314	43682	A	13394	3119	6984	CIVKHAAMRSAMTSLAARITVL NPTRATTPNT/RVADDQGFLRQ WSKVAKERKLQRLYIGEPSAEA
Ì					·	VAAQMPDLILISATGGDSALAL
						YDQLSTIAPTLIINYDDKSWHPR
į						LSFNKADEITTVFCGSKKSLAN
j						GIPMANILFPTSVIATQQTQHFR
}		ŀ				FVDDFNAQFLRFASLDPAASPA
						ITISGKATLEIPGTELVFYAALE
						KKLNIKPGQTTFDGRFTLLPTC
		-				CLGNCDKGPNMMIDEDTHAHL
						TPEAIPELLER
13315	43683	Α	13395	3	587	
13316	43684	Α	13396	1	315	
13317	43685	A	13397	430	636	
13318	43686	Α	13398	1	84	
13319	43687	Α	13399	l	1647	
13320	43688	A	13400	1	2652	
13321	43689	В	13401	1	1494	
13322	43690	Α	13402	1	750	
13323	43691	Α	13403	1	513	
13324	43692	Α	13404	1	2254	
13325	43693	Α	13405	1	746	
13326	43694	Α	13406	3	428	LGDDFVRAANIILHCEGKVVVS
	1					GIGKSGHIGKKIAATLASTGTPA
						FFVHPAEALHGDLGMIESRDV
						MLFISYSGGAKELDLIIPRLEDK
						SIAL\GKPTSPLGLAAKAVLDIS
						VEREACPMHLAPTSSTVNTLM
		<u> </u>		-		MGDALAMAV
13327	43695	Α	13407	1	1431	
13328	43696	A	13408	943	2481	
13329	43697	В	13409	1	783	
13330	43698	A	13410	3	729	
13331	43699	A	13411	1	843	
13332	43700	A	13412	3	128	
13333	43701	A	13413	1	136	
13334	43702	A	13414	1	492	
13335	43703	Α	13415	3	619	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
13336	43704	Α	13416	1	1600	GQACHASSSPLKGSRGSPPNFEI
						LYGPIFEDSLAPD*PKVIAGQ\IT
						LRDAVTGPISSPNEAGKIYQLKP
						NPAVLICRVRGLHLPEKHVTW
						RGEAIPGSLFDFALYFFHNYQA
						LLAKGSGPYFYLPKTQSWQEA
						AWWSEVFSYAEDRFNLPRGTIK
						ATLLIETLPAVFQMDEILHALR
						DHIVGLNCGRWDYIFSYIKTLK
						NYPDRVLPDRQAVTMDKPFLN
				ł		AYSRLLIKTCHKRGAFAMGGM
						AAFIPSKDEEHNNQVLNKVKA
						DKSLEANNGHDGTWIAHPGLA
						DTAMAVFNDILGSRKNQLEVM
		1				REQHAPITADQLLAPCDGERTE
ŀ						RKFNALMESEKGQSQNLWKFA
						VYSGLRHGELAALAWEDVDLE
						KGIVNVRRNLTILDMFGPPKTN
						AGIRTVTLLQPALEALKEQYKL
						TGHHRKSEITFYHREYGRTEKQ
						KLHFVFMPRVCNGKQKPYYSV
						SSLGARWNAAVKRAGIRRRNP
						YHTRHTFACWLLTAGANPAFIA
		ŀ				SQMGHETAQMVYEIYGMWID
						DMSDEQIAMLNARVM
13337	43705	Α	13417	2	278	
13338	43706	Α	13418	3	662	GHLWIRIDLSQSAVSHSVKELE
						NHTGVRLLDRTTREVVLTDAG
1						QQLALRLERLLDELNSTLRDTG
		Ì				RMGQQLSGKVRVAASQTISAH
						LIPQCIAESHRRYPDIQFVLHDR
						PQQWVMESIRQGDVDFGIVIDP
			ŀ			GPVGDLQCEAILSEPFFLLCHRD
						SALAVEDYVP\ALPLPEGSPLVV
						KRITPVVERQLMLVRRKNRSLS
			ļ. <u> </u>			TAAEALWDVVRDQGNALMAA
13339	43707	Α	13419	1	646	GQQLSGK\VRVAASQTISAHLIP
						QCIAESHRRYPDIQFVLHDRPQ
1						QWVMESIRQGDVDFGIVIDPGP
						VGDLQCEAILSEPFFLLCHRDSA
					·	LAGGDYVPWQALQGAKLVLQ
						DYASGSRPLIDAALARNGIQAN
						IVQEIGHPATLFPMVAAGIGISIL
						PALALPLPEGSPLVVKRITPVVE
						RQLMLVRRKNRSLSTAAEALW
		_				DVVRDQGNALMA\GRE
13340	43708	Α	13420	1	1068	
13341	43709	A	13421	1	381	
13342	43710	A	13422	3	938	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
		<u> </u>				
13343	43711	Α	13423	595	1630	CSWHDRFPDWKAGRILPISEPPS
						NRIFACWGKPAWTACCN/FSQG
						QAVKGNQLLPVSLVKRKTTLA
			:			PNTQTASPRALADSLMQLARQ
						VSRLESGQDFADFGTTIKQDFR
					İ	LLGQTSVDRLLQLSQGQAVKG
			•		1	NQLLPVSLVKRKTTLAPNTQTA
		l				SPRALADSLMQLARQVSRLESG
					ł	QDFADFGTTIKQDFRLLGQTSV
						DRLLQLSQGQAVKGNQLLPVS
						LVKRKTTLAPNTQTASPRALAD
						SLMQLARQVSRLESGQDFADF
						GTTIKQDFRLLGQTSVDRLLQL
			•			SQGQAVKGNQLLPVSLVKRKT
			[TLAPNTQTASPRALADSLMQLA
1						RQVSRLESGHYDHDYEFELGTR
						FRAVIIKLCNGDSLRHFMTSVY
						ALVKCFHEGDPLHKTGCQHLIP
						KQQQNEKYQVPQFDQSTIKNIE
						SAKGLDVWDSWPLQNADGTV
						AEYNGYHVEMLLREAGKAGFC
						RFRNHHQTGFSPAGANQRGPL
						AATLSGPGGEGQSAVARLTGE
						KKNHPGAQYANRLSPRVGRFIN
						AAGTTGNSYDHDYEFELGTHQ
						RLGAVARIVGDDLLNELRRDV
						VDARAGDVDQHAHAEKGDFR
13344	43712	Α	13424	1	1395	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13345	43713	A	13425	1307	1992	DKLNPVAHQKAYLAQSSQCHP QNARLVQHTQIRILPISEPPSNRI FACWGKPAWTACCN/FSQGQA VKGNQLLPVSLVKRKTTLAPNT QTASPRALADSLMQLARQVSR LESGQDFADFGTTIKQDFRLLG QTSVDRLLQLSQGQAVKGNQL LPVSLVKRKTTLAPNTQTASPR ALADSLMQLARQVSRLESGQD FADFGTTIKQDFRLLGQTSVDR LLQLSQGQAVKGNQLLPVSLV KRKTTLAPNTQTASPRALADSL MQLARQVSRLESGQDFADFGT TIKQDFRLLGQTSVDRLLQLSQ GQAVKGNQLLPVSLVKRKTTL APNTQTASPRALADSLMQLAR
						QVSRLESGQLKHKEVESPNRPI TSSEIQVLIKSLPIKKSPGPDGFT AEFYERYKEELVPFLLKLFQTIE QERLLPNSFYEASIILTSKPSRD ATKKDNFRPTSLMNIYAKILNK IGQTESSSTSKSLFSTIKSVSSPE CKAGSTYANQDFADFGTTIKQ DFRLLGQTSVDRLLQLSQGQA VKGNQLLPVSLVKRKTTLAPNT QTASPRALADSLMQLARQVSR LESGQETTPWSPFSDPVLAFDY KVFMAFVYNVVRVSQRMVVA GCISAFNDKLLNDLICFPVAPVS SKQIFPSDVKVAEPDHSCVSSY RIGSMAPNLTLSEMPRHNYFPA GNSVDHDNEFEL
13346	43714	A	13426	560	844	
13347 13348	43715	B A	13427 13428	950	1519 1230	CSWHDRFPDWKAGRILPISEPPS NRIFACWGKPAWTACCN/FSQG QAVKGNQLLPVSLVKRKTTLA PNTQTASPRALADSLMQLARQ VSRLESGQDFADFGTTIKQDFR LLGQTSVDRLLQLSQGQAVKG NQLLPVSLVKRKTTLAPNTQTA SPRALADSLMQLARQVSRLESG QDFADFGTTIKQDFRLLGQTSV DRLLQLSQGQAVKGNQLLPVS LVKRKTTLAPNTQTASPRALAD SLMQLARQVSRLESGQDFADF GTTIKQDFRLLGQTSVDRLLQL SQGQAVKGNQLLPVSLVKRKT TLAPNTQTASPRALADSLMQLA RQVSRLESGQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13349	43717	A	13429	695	1060	FRQRSVAFPPTVPNACVSVRSR ERDFADFGTTIKQDFRLLGQTS VDRLLQLSQGQAVKGNQLLPV SLVKRKTTLAPNTQTASPRALA DSLMQLARQVSRLESGQ*AQR N*CRKHRLSIPRK
13350	43718	A	13430		1558	MRDIFWVPDICKVMLYLVFFRS MGTPLRAVAMGLVITSWKIAQ KPRDFADFGTTINQDFRLLGQT SVDRLLQLSQGQAVKGNQLLP VSLKTDTKAKNLYLTKAYYGG GTNFFRKESQKLQQSAKKRDA ELANGALGIIELNNDYTLKKVM KPLITSNTGFADFGTTIKQDFRL LGQTSVDRLLQLSQGQAVKGN QLLPVSLVKRKTTLAPNTQTAS PRALADSLMQLARQVSRLESG Q*AQRN
13351	43719	A	13431	391	1676	FIRDFADFGTTIKQDFRLLGQTS VDRLLQLSQGQAVKGNQLLPV SLVKRKTTLAPNTQTASPRALA DSLMQLARQVSRLESGQ*AQR N
13352	43720	В	13432	1	1638	
13353	43721	A	13433	1	2430	
13354	43722	В	13434	1	1282	
13355	43723	В	13435	1	1225	
13356	43724	Α	13436	350	709	
13357	43725	A	13437	548	2236 1698	RSWLRSSLSGSIPIATEISCTSCE ASTSS/VRDFPGVKDLTFQWHD RLIFAIARLLGRSACGVTFHKEQ LGTVKVLRGTISQFARQRRAAG QLFTHHFFGRTHTALGAGNRH LRQQFSRLNVLVQPQDKGIFHH ARNECRALTRRETPAFFRIRAVF NSQLSQTQLHIFFTARPAGLAS KCRKPPQACYLWDFADFGTTIK QDFRLLGQTSVDRLLQLSQGQ AVKGNQLLPVSLVKRKTTLAP NTQTASPRALADSLMQLARQV SRLESGQNIESAKGLHVWDSW PLHNADGTVDEYNGYHVVFAL AGSPKDADDTSIYMFYQKVGD NSIDSWKNAGRVFKDSDKFDA NDPILKDQTQEWSGSATFTSDG KIRLFYTDYSGEHYG
13359	43727	В	13439	448	1116	
13360	43728	A	13440	97	1636	
13361	43729	A	13441	1	1296	
13362	43730	В	13442	1	2160	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	ŀ	09/340,417	sequence	of peptide sequence	defetion, (-possible nucleotide insertion)
13363	43731	Α	13443	485	865	FIRDFADFGTTIKQDFRLLGQTS
						VDRLLQLSQGQAVKGNQLLPV
		ļ				SLVKRKTTLAPNTQTASPRALA
						DSLMQLARQVSRLESGQS*NSS
						KTKNTKCLNSINQRLKILSLQK
	}					DLMCGTAGRCKTLTEQ
13364	43732	A	13444	816	1194	SLILFLRRRRAVEKRGKVWKDS
						CPNGTTLTLSGAIFLFIRDFADF
		l				GTPITQDFRLLGQTSVDRLLHLS
						QGQAVKGNQLLPVSLVKRKTT
1			ļ			LAPNTQTASPRALADSLMQLA
ŀ						RQVSRLESGQ*AQRN
13365	43733	В	13445	İ	2710	
13366	43734	A	13446	2027	2934	NPRILPISEPPSNRIFACWGKPA
						WTACCN/FSQGQAVKGNQLLP
		ŀ				VSLVKRKTTLAPNTQTASPRAL
		l				ADSLMQLARQVSRLESGQDFA
						DFGTTIKQDFRLLGQTSVDRLL
						QLSQGQAVKGNQLLPVSLVKR
						KTTLAPNTQTASPRALADSLMQ
	,					LARQVSRLESGQSRVHSHSLGP
						LPTTTTGSLINQKKGGVSGGPG
		ŀ				SKEQAEISCLKERKDLGLGILGS
		·				TVGILKVKHQGASTKQGHSQD
		ļ				LIKILLSDTSLSTPAAPMVDSLIA
		1				RVGVMARGNAITLPVCGRDVK
						FTLEVLRGDSVEKTSRVWSGNE
						RDQELLTEDALDDLIPSFLLTGQ
						QTPAFGRRVSGVIEIADGSRRR
						KAAALTESDYRVLVGELDDEQ
						MAALSRLGNDYRPTSAYERGQ
						RYASRLQYMNLLPIFLRWLMR
						KIFHDEHGVPQDQQEEEPISETII
		İ				EEEIPSINSDLGNELYFVKLPKF
						LSIEPKDFADFGTTIKQDFRLLG
						QTSVDRLLQLSQGQAVKGNQL
						LPVSLVKRKTTLAPNTQTASPR
						ALADSLMQLARQVSRLESGQP
ł						HSTDSATHRKMTLSLADRCSKT
						QKIRILPMAGRDPECQRTEMIK
						SSDLMAEIIQEHKEDQLPELEQL
					†	EHIGLFSHAEIKAIIKKASDLQY
						RIQERALFKDDFINYVLHEIPLF
						EQIQRRTRIGYSLKHGIRAVKA
	I	ļ				TIVRALYGRIKPAGVVVTRSVP
13367	43735	A	13447	I I	5240	
13368	43736	A	13448	4733	4753	

SEQ ID NO:	SEQ ID NO: of peptide	I	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	liou	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13369	43737	Α	13449	299	761	LNTTAITLCLLLREARKTLMTH
						QSTCFIKRDFADFGTTIKQDFRL
						LGQTSVDRLLQLSQGQAVKGN
						QLLPVSLVKRKTTLAPNTQTAS
						PRALADSLMQLARQVSRLESG
						Q*AQRN**PVPKEKRRRGDRQE
					44.00	HQQPCRNTEGTALRVWAARDQ
13370	43738	A	13450	3925	4190	FIRDFADFGTTIKQDFRLLGQTS
						VDRLLQLSQGQAVKGNQLLPV
	ĺ					SLVKRKTTLAPNTQTASPRALA
						DSLMQLARQVSRLESGQ*AQR
13371	43739	A	13451	2	1040	N
13371	43740	A	13451	58	350	HLASWVVALVLHWGCVWVTA
13372	43740	^	13432	36	1330	ESSTSKTRSPFRARSWIGEAVA
						AAAADGVTFSVPVTPHTFRHSY
						AMHMLYAGIPLKVL\QSLMGH
						KSISSTGVPSSNS
13373	43741	A	13453	1	1290	
13374	43742	A	13454	289	1338	
13375	43743	Α	13455	ı	732	
13376	43744	A	13456	982	1347	
13377	43745	A	13457	116	1746	SSVTGRTEKARIWEVTDRTV/R
						RPWIGEAVAAAAADGVTFSVP
		ĺ				VTPHTFRHSYAMHMLYAGIPL
						KVLQSLMGHKSISSTEVYTKVF
						ALDVAARHRNRFTQFRLSETKE
						ITNPYAMRLYESLCQYRKPDGS
						CIVSLKIDWIIERYQLPQSYQLY
						YFELAIPVGYFYPGSFSTASRIL
ŀ				:		LLHPRGLRAITIAVFGKQNTYIR
						LEPFKINVLEQITKHIEKLQCGG
						VVKQLSRRGNNQHISSTYDINR ADTQVRRAVNNYDIIVMSNSFN
						GQSEHQVWIARLTWVIGTINVV
						CAADVLIVPTPAELFDYTSALQ
						FFDMLRDLLKNVDLKGFEPDV
						RILLTKYSNSNGSQSPWMEEQI
						RDAWGSMVLKNVVRETDEVG
						KARLTWGIGTINVVCAADVLIV
1						PTPAELFDYTSALQFFDMLRDL
						LKNVDLKGFEPDVRILLTKYSN
						SNGSQSPWMEEQIRDAWGSMV
						LKNVVRETDEVGKEPPSTNTFR
						HSYAMHMLYAGIPLKVLQSLM
						GHKSISSTEVYTKVFALDVAAR
						HRVQFAMPESDAVAMLKQLS
13378	43746	Α	13458	1	780	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13379	43747	A	13459	116	396	SSVTGRTEKARIWEVTDRTV/R RPWIGEAVAAAAADGVTFSVP VTPHTFRHSYAMHMLYAGIPL KVLQSLMGHKSISSTEVYTKVF ALDVAARHR
13380	43748	Α	13460	3	514	
13381	43749	A	13461	2	545	
13382	43750	Α	13462	1	1110	
13383	43751	A	13463	1	870	
13384	43752	A	13464	1	1422	
13385	43753	Α	13465	1	1599	
13386	43754	A	13466	614	1236	TARVAAARRRCQCWGSGACGS ALTLPRTSSLTLTETPIVQQTRM VVRCPTASLPSQWRSCPSSTWA VPKTPPTWTCWRNSAILSL/VVF IRDKLMERRNRRTGRTEKARIW EVTDRTVRTWIGEAVAAAAAD GVTFSVPVTPHTFRHSYAMHM LYAGIPLKVLQSLMGHKSISSTE VYTKVFALDVAARHRVQFAMP ESDAVAMLKQLS
13387	43755	Α	13467	1	1023	
13388	43756	Α	13468	1	699	
13389	43757	В	13469	1	1281	
13390	43758	В	13470	77	751	
13391	43759	Α	13471	1	705	
13392	43760	Α	13472	2	893	
13393	43761	A	13473	206	837	CAGAGGIARRRAEKEKALSAV NLDPTSRRLSSTSLPGAPPNPRV AAAGLSRRRPTPESAGRKS/GR WPRASRYQQGHQDLFILRSDLP SQVFIRDKLMERRNRRTGRTEK ARIWEVTDRTVRTWIGEAVAA AAADGVTFSVPVTPHTFRHSYA MHMLYAGIPLKVLQSLMGHKS ISSTEVYTKVFALDVAARHRVQ FAMPESDAVAMLKQLS
13394	43762	A	13474	1	1416	
13395	43763	A	13475	1	765	
13396	43764	В	13476	1	519	
13397	43765	A	13477	1	2784	
13398	43766	A	13478	1	624	
13399	43767	Α	13479	2089	2454	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13400	43768	Α	13480	740	1431	CCQTLPVFHPPHPSHLGPRPCPP
			·		•	TH/HLPSEHRSLFLEAACHDSDS
						LEPLNLSSGSKTKSPSLPPKAKK
						PKGLEISA/RPAGALRHRHRLHR
						PQQPSPPLGIPHPSLL\TAQVFIR
						DKLMERRNRRTGRTEKARIWE
				1		VTDRTVRTWIGEAVAAAAADG
						VTFSVPVTPHTFRHSYAMHML
						YAGIPLKVLQSLMGHKSISSTE
		1				VYTKVFALDVAARHRVQFAMP
		1				ESDAVAMLKQLS
13401	43769	Α	13481	2	2309	
13402	43770	Α	13482	1	603	
13403	43771	Α	13483	1	762	MVPHSTGKSWNGCDPTSATLD
		1				QLVTVFFMKLFSIGDARIPCLGP
						RSHSYRRSDYYSGTTIHSCCRA
		1				DYWSGTTAHLYCRSDYHPVPS
				l		FILVTSTTNHQPLKPSISEASWN
						PLFASL/HHLQLITES/W/YMSLA
						TTVFIRDKLMERRNRRTGRTEK
		1		1		ARIWEVTDRTVRTWIGEAVAA
						AAADGVTFSVPVTPHTFRHSYA
						MHMLYAGIPLKVLQSLMGHKS
						ISSTEVYTKVFALDVAARHRVQ
						FAMPESDAVAMLKQLS
13404	43772	A	13484	1	762	
13405	43773	A	13485	1	615	DDCV + D (HVCV/TDDTV/DTV/DD
13406	43774	Α	13486	887	1203	RPEKAR/IWGVTDRTVRTWIGR
						AVAAAAADGVTFSVPVPPHTF
						RHSYAMHMLYAGIPLKVLQSL
						MGHKSISSTEVYTKVFALDVAA
12407	12775		12407	<u> </u>	055	RHRVQFAMPESDAVAMLKQLS
13407 13408	43775	A	13487 13488	1	855 927	
13408	43777	A	13489	1	921	
13410	43778	A	13490	1	3216	
13411	43779	A	13491	1	1974	
13411	43779	A	13491	1	1221	
13412	43781	В	13493	1	2337	
13413	43782	A	13494	1	1273	
13414	43783	A	13494	1	1251	
13415	43784	A	13496	1999	3237	
13417	43785	A	13490	1	1773	
1341/	43/83	A	1349/]1	11/13	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13418	43786	A	13498		633	AQALIATYGRDRDPGRPLWLG
13	13700	` `	13.75			SVKSNIGHTQAAAGVAGVIKM
						VMAMRHGQLPRTLHVESPSPE
						VFIRDKLMERRNRRTGRTEKAR
			ŀ			IWEVTDRTVRTWIGEAVAAAA
						ADGVTFSVPVTPHTFRFSYAMH
1			1	-		MLYAGIPL*VLQSLMGHKSISS
		İ				TEVYTKVFALDVAARHRVQFA
		1				MPESDAGGPGTCACSQQGCRL
						RGSSVSPGRQHPQPHVRV
13419	43787	A	13499	101	1943	
13420	43788	Α	13500	686	3925	
13421	43789	Α	13501	1	2084	
13422	43790	Α	13502	734	834	
13423	43791	Α	13503	1	1075	
13424	43792	Α	13504	1	357	
13425	43793	Α	13505	795	1273	SHPFFSTDWSTDTGFRSKSIWC
						HRNCRW/RVAVVKLLHLPKVII
						VFWLASWMMSRWLHYPDWVT
		İ				IIAQQVLMNVVSVMQADCRMN
			<u> </u>			LLEIFLRWLMRKIFHVRLLPAV
						STPPNCLNQLLLFFLTPVNYLPG
						QVMHFKKPLQIKRNYLSSRHLT
		1				FMSRKKLG
13426	43794	Α	13506	344	449	SHPFFSTDWSTDTGVRSKSIWC
		ı				HRNCRW/RVAVVKLLHLPKVII
						VFWLASCMMSRCPHYPDWVTI
		l				IAQQVLTPRKDVMQADCRMNL
						LEIFLRWLMRKIFHGFGM
13427	43795	Α	13507	1197	1511	WHYREPLLLRAQLTNPPSGQLV
				İ		SITKYLQPVEY*AYVLE*QAHL
ł					[HYQLKSVDQWPVQKQFSSGSD
						LYQLHPFHVQHFLEPCFPRHPE
						FAPPSTGTESHYYCCIW
13428	43796	В	13508	1	1494	
13429	43797	Α	13509	1	554	MTARKARKITRRWRIGEAADL
						VGVSSQAIRDAEKAGRLPHPD
						MEIRGRVEQRVGYTIEQINHMR
						DVFGTRLRRAEDVFPPVIGVAA
				}		HKGNDPQGTASMYHGWVPDL
						HIHAEDTLLPFYLGEKDDVTYA
						IKPTCWPGLDIIPSCLALHRIETE
						LMGKFDEAQPNLGIGT/N*CRM
				<u> </u>		CC*CADCSHAC
13430	43798	Α	13510	1771	2068	DTYSVSWIGEAVAAAAADGVT
				1		FSVPVTPHTFRHSYAMHMLYA
		1				GIPLKVLQSQYPIFKLSYKNPVT
						QTAWF\WFPNRIFGPVPINVSFS
	<u> </u>					PKVGPGLPGII
13431	43799	Α	13511	65	277	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13432	43800	A	13512	327	748	VHQFHGSLHEGFCAGCGCPAP
						GTVCDAGGFCRFRNHHQTGFSP
						AGANQRGPLAATLSGPGGEGQ
		1				SAVARLTGEKKNHPGAQYANR
		1				LSPRVGRFINAAGTTGFPTGKR
		ŀ				AKSIWCHRNCRW/RVAVVKLL
						HLPKVIIVFWLAS
13433	43801	Α	13513	512	580	
13434	43802	A	13514	2096	2272	EWRNVCGVTGTENVTPSAAAA
		ŀ	į			QPPPQSRPFSWILRPF*RP*RKIS
						TSFIRPLFTFLPA
13435	43803	Α	13515	402	2784	RKNPFILH*LFR*TLRQTKPDNS
						AGKCVKI**HTQNQRSGRSQSD
						F*RRRKNISERSAAYR*RQLYIR
						RQPYAERPS\HVEDKGHKYLVF
						EANTGTENGYQGEESLFNKAY
		ł				YGGGTNFFRKESQKLQQSAKK
					}	RDAELANGALGIIELNNDYTLK
						KVMKPLITSNTVTDEIERANVF
						KMNGKWYLFTDSRGSKMTIDG
						INSNDIYMLGYVSNSLTGPYKP
			ŀ			LNKTGLVLQMGLDPNDVTFTY
						SHFAVPQAKGNNVVITSYMTN
						RGFFEDKKATFAPSFLMNIKGN
						KTSVVKNSILEQGQLTWLQVA
					1	KRAGLGGGQSGRTVLRERLPN
						YKNFKGTIQELGQNQYAVSGEI
		ŀ				FVVDRNTVEITELPVRTWTQVY
		l				KEQVLEPMLNGTDKTPALISDY
						KEYHTDTTVKFVVKMTEEKLA
						QAEAAGLHKVFKLQTTLTCNS
						MVLFDHMGCLKKYETVQDILK
						EFFDLRLSYYGLRKEWLVGML
						GAESTKLNNQARFILEKIQGKIT
		ŀ				IENRSKKDLIQMLVQRGYESDP
		l				VKAWKEAQEKCDNLKTCHTSH
		l	ļ			GSVMAETAVINHKKRKNSPRIV
						QSNDLTEAAYSLSRDQKRMLY
						LFVDQIRKSDGTLQEHDGICEIH
						VAKYAEIFGLTSAEASKDIRQA
						LKSFAGKEVVFYRPEEDAGDE
						KGYESFPWFIKRAHSPSRGLYS
						VHINPYLIPFFIGLQNRFTQFRLS
13436	43804	Α	13516	2	199	WRKKSLDIPPLIIPNGIVKNILRH
		1				FSQLLNVPITRPFSWILRPF*RP*
		L				RKISTSFIRPLFTFLPA
13437	43805	Α	13517	1	1887	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13438	43806	Α	13518	125	828	ECRKHTKLKLQPTIWHVLKI/Y
<u> </u>						NSEKLLVYRKLLQVLTGLYKRS
		ŀ				NLGVRLYACCGLLLCPAYPQH
						FAHGYVDKIPGYPGRAGTLTGL
						HPMQCDNLKTCHTSHGSVMAE
	:					TAVINHKKRKNSPRIVQSNDLT
						EAAYSLSRDQKRMLYLFVDQIR
:						KSDGTLQEHDGICEIHVAKYAE
						IFGLTSAEAILRYFIKHSADMEA
						ITNGMMNLNRQRHQHLVALRII
						FAHGENGGEEVVHIGHV
13439	43807	Α	13519	504	755	EPCFPAHPEFAPLSTGAESHHY
13439	43607	<u> </u>	13313	304	1733	CCIW*AKYVHQARTL*DQRS*A
		ĺ				DHEAYRKTAVRRCSQTTQQAW
12440	42000	n	12520	1	1000	EQLVHQQHILPSSCRYPG
13440	43808	В	13520	1000	1980	
13441	43809	A	13521	923	1063	-
13442	43810	В	13522	1	1932	DEVERTAL EDMONARE AND E
13443	43811	A	13523	383	1383	RKVFFIALKRMPAMKKAMNLF
						LGLSNVRTVHPEGFTVYISTHIS
		l				FPSLSGYKTRLRSFGLVKQKKS
		1				PILLEVLAREIRQE/KEIKGIQFG
		ļ		1		KEEVKLSLFADDMIVYLENPIV
						SAQNLLKLISNFSKVSGYKINV
		ĺ				QKSQAFLYTNNRQTESQIMSEL
					•	PFTIASKRIKYLGIQLTRDVKDL
						FKENYKPLLNEIKEDTNKWKNI
						PCSWVGRINIVKMAILPKLSRM
						HGRPHQPPNAASTLPFPKRRSH
						YGQKSERLSRSGDNRKSHPECR
	1					HWQSCAKKSRCLAFTRTVRHT
						PRSAYLLTRTIELLESSALWHV
						QTQRIYAPLYETLDPGNRTHCE
13444	43812	В	13524	107	1000	
13445	43813	A	13525	1778	1897	
13446	43814	Α	13526	273	421	SHPFFSTDWSTDTGVRSKSIWC
						HRNCRW/RVAVVKLLHLPKVII
	<u> </u>	L				VFSSAS
13447	43815	Α	13527	273	697	SHPFFSTDWSTDTGVRSKSIWC
		l				HRNCRW/RVAVVKLLHLPKVII
						VFWLASWMMSRWLHYPDWVT
				1		IIAQQVLMNVVSVMQADCRMN
						LLEIFLRWLMRIIFHVRLLPAVS
						TPTNCLNQLLLFFLTPVNYLPG
						QSGTTVPLVSSV
L	<u> </u>	<u> </u>		L	<u> </u>	G9011ALTA99A

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
:				sequence		
13448	43816	Α	13528	273	688	SHPFFSTDWSTDTGVRSKSIWC
						HRNCRW/RVAVVKLLHLPKVII
						VFWLASWMMSRWLHYPDWVT
1						IIAQQVLMNVGSVMPADCRMN
ŀ		Ī				LLEIFLRWLMRKIFTTGLRSFGL
						VKQKKSPIRMPCVYTNPCVSIV
						SRMGQASSL
13449	43817	В	13529	1	1053	old violet
13450	43818	В	13530	1	1869	
13451	43819	A	13531	120	430	EDLSDTGVRSKSIWCHRNCRW/
13431	43017	Α	15551	120	1750	RVAVVKLLHLPKVIIVFWLASW
				:		MMSRWLHYPDWVTIIAQQVL
1						MNVVSVMQADCRMNLLEIFLR
1						WLMRKIFHSSLEDPRNWHL
13452	43820	В	13532	148	334	WEMKRIFHSSLEDFKNWHL
13452	43821	A	13532	2	147	TSMDKGLQGALKKKI*DL*KK*
13433	43821	l ^A	13333	2	147	L\KNIPVDKWITPREVFSGIVCK
						ONI
13454	43822		13534	151	325	KAICHNTFSKPSMDKGL/LRRS
13434	43822	Α	13334	151	323	l l
						KKEGFKTYKKNIYKNIPVDKWI
12455	42022	_	12525	202	406	TPREVFSGIVCKQNI
13455	43823	Α	13535	283	406	HRTKISIVPEGIFKSFSLPLAGIM
10456	1000.1	_	10506		1000	TEQYFL*TLHRLLRLIM
13456	43824	В	13536	1	1902	L HIM E A CIVIDIDO A LIVOS A CIVIDIV
13457	43825	A	13537	508	1041	LWREASKDIRQALKSFAGKEV
						VFYRPEEDAGDEKGYESFPW/C
		l				YQT\WHSPSRGLYSVHINPYLIP
						FFIGLQNRFTQFRLSETKEITNP
						YAMRLYESLCQYRKPDGSGIVS
						LKIDWIIERYQLPQSYQRMPDF
						RRRFLQVCVNEINSRTPMRLSYI
						EKKKGRQTTHIVFSFRDITSMTT
13458	43826	Α	13538	1	1422	
13459	43827	В	13539	1	1002	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13460	43828	Α	13540	1	160	MVLIVAATVVVVVVVGLCWM
	:	1				AKSECRSAMRLSSLQVRIFTYLI
						GREAAFADNLKWMACANKES
						ALLDRFIFLILLLCKEPWECHCL
						PLFKVAMPSKCDIPVICAHINIG
					i i	FTLRQLRGFGPCNIGSFEAKDS
						KNPQVRVISQKHKALRCIWFIT
						CSICNPEFTEGALQGNKQRKRQ
						CLQWFRAKALESNEPGFEFIQD
						KLLTQSLCFFPYTFLWLTGPHPS
		ļ				GSSYTDPCSSKADVTPSSTAVQ
						EHSSALELKTKIVSPITRQQHGF
						FTQISTLADVQENVMEYLHVLS
	1	ŀ				RPKVIDQEHDVVWTEAYIDSTV
						CRVKRRQLLLQQVPLTRKQQF
						LPAPYVDLDSARTRTNKQQQN
						PIIYTGSKQGSHVILFHYGESWN
		ŀ				LLRADQRLIFAKSWPRASRYQQ
						GHQDLFILRSDLPSQVFIRDKL
		l				MERRNRRTGRTEKARIWEVTD
						RTVRTWIGEAVAAAAADGVTF
		l				SVPVTPHTFRHSYAMHMLYAG
						IPLKVLQSLMGHKSISSTEVYTK
					i e	VFALDVAARHRVQFAMPESDA
						VAMLKQL*QPQWWWWSW*D
						CAGWRKVNAAQPCVCLPFRFA
12461	43829		13541	40	668	SSHTSLDERLRLQTI
13461 13462	43829	A	13542	48	1278	
13463	43831	A	13543	100	157	
13464	43832	A	13544	1	1122	
13465	43833	В	13545	1	1131	
13466	43834	Ā	13546	273	812	SHPFFSTDWSTDTGVRSKSIWC
		-				HRNCRWESPS*SC/WHLPKVIIV
						FWLASWMMSRWLNYPDWVTII
						AQQVLMNVVSVMQADC\KRIC
						WKYFCAG*CGKIFTYQ**TW\N
						NTETSMSQKPCKMKRI/YSREE
		1				LAV*KMDHHSDLISLW*KLEPL
						TCRSTSHFRQKLAQGFPVSTGT
						PGFIYSAK
13467	43835	В	13547	1	987	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13468	43836	A	13548	246	2065	LVGWLDGWLDGWLIVALPFM MIPWHSSIQPFTIHPFNWNFG*M PELCPQI\GMDSWLAGWM/CWL DGWLDGW*MAKLRKPVL*PD KEGNEIWVDMYTVKPSGWTVR TFDKPRKRFIAFFIAGILFRAIKN HFLPRETLQCLPYILTGFRRGQS EYFSIFSNMDLADTVMFLIVAL PFMMIPWHSSIQPFTIHPFNWNF G
13469	43837	Α	13549	2	118	
13471	43838	A	13550	147	417	THGIRIGDFFCFTKRKLRKPVL* PDKEGNEIWVDMYTVKPSGWT VRTFDKPRKRFIAFFIAGILFRAI KNHFLPRETLQCLPYILTGFRRG QSEYFSIFSNMDLADTVMFL SSSWGRSSPQLVCGIFQTAGPQS LCKWNQ*VLAQTNKKTAGTSA DLYVPV*QL*RKAVVLPTNAP WRFWRTRQEWFSSRGRSGYQS QET
13472	43840	A	13552	1	2418	<u> </u>
13473	43841	В	13553	1	1848	
13474	43842	A	13554	188	490	EWRNVCGVTGTEKVTPSAAAA ATASPIQVLTVLSVTSQIRAFSV LPVRRLRRSMSLSRINTCVNRF CNPIKKGMRYGLICPL*SLLDGL CARLINQGKDS
13475	43843	A	13555	83	440	SLLFKCSGVIVLRRPLGYRQVM NVTRRCLLRTHWMISSLLFY*L VNRHRRSVEEYLVSYTLPMGV AVVKLLHLPKNFVSLITPRGRL EKERAFLFQHMRSISTGLKIFV MHTSCPMVVC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		<u> </u>				
13476	43844	Α	13556	1337	2583	QQRR*MCQNLMTH\QNQRSGR
				:		SQNDF*RRRKNISERSAVYR*R
						QLYIRRQPYAERPS\HVEDKGH
		1				KYLVFEANTGTENGYQGEESLF
						NKAYYGGGTNFFRKESQKLQQ
		1				SAKKRDAELANGALGIIELNND
						YTLKKVMKPLITSNTVFIRDKL
		1				MERRNRRTGRTEKARIWEVTD
						RTVRTWIGEAVAAAAADGVTF
		l				SVPVTPHTFRHSYAMHMLYAG
						IPLKVLQSLMGHKSISSTEVYTK
ļ						VFALDVAARHRIRKSDGTLQEH
ŀ						DGICEIHVAKYAEIFGLTSAEAS
		İ				KDIRQALKSFAGKEVVFYRPEE
						DAGDEKGYESFPWFIKRAHSPS
						RGLYSVHINPYLIPFFIGLQNRF
						TOFRLSETKEITNPYAMRVTNP
		ł				CVSIVCVNEINSRTPMRLSYIEK
		1		İ		KKGRQTTHIVFSFRDITSMTTG
13477	43845	В	13557	1	696	
13478	43846	A	13558	326	1577	SGLLPIKVAFTKPQFLFILLRIW*
						AKYVHQARTL*DQRF*ADHEA
•						YRKTAVRRCSQTTQQAWEQSA
•						HQQHIRH
13479	43847	В	13559	1	3822	
13480	43848	Α	13560	848	1476	VNVTSFGSRPICSTSPVLFSGL*
						GPVKEFDT*PSM*ISFELIPSIVIF
						EPRESVNKYHLPFILKTFARSISS
						VTVFEVISGFITFFNV*SLFNSMI
Į						PRAPFANSASRFLALC*SFWLSL
						RKKFVPPP*YALLNKDSSPW/SS
		ı				VFCSRVGFEYKVFMAFVFNVV
						RVSQRMVVAGCIIAFIDKLLNV
				<u></u>		LICFSVSVKNRFVIFHSVDFECV
13481	43849	Α	13561	1	2196	
13482	43850	Α	13562	326	3746	SGLLPIKVAFTKPQFLFILLRIW*
						AKYVHQARTL*DQRS*ADHEA
				-		YRKTAVRRCSQTTQQAWEQSA
						HQQHIRH*SCRYPS*ARCQ*LTG
						QIVHRGEKKEQQLI
13483	43851	Α	13563	3802	4075	IRCFTYSPAPGYFPRFRAVIIKLC
						NGDSLRHFMTSVYALQEAAAE
						VRHTLVTLRQLVTLYDPVDFQ
						RDDA*AIRAYDTDTGIRINAWH
						TDW
13484	43852	Α	13564	1708	3198	
13485	43853	A	13565	1	1401	NAME OF THE PARTY
13486	43854	Α	13566	1214	1357	NKINMFIAALFTIAKT\WNQPK\
		1				CPTMIDWIKKRGSSRVASSSSPT
	<u> </u>	1				RTR

1	of peptide sequence	hođ	in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13487	43855	A	13567	3363	3852	SHPFFSTDWSSDTGVRSKSIWC HRNCRW/RVAVVKLLHLPKVII VFWLASWMMSRWLHYPDWVI IIAQQVLMNVVSVMQADCRMN LLEIFLRWLMRKIFHVRLLPAV STPPNCLNQLLLFFLTPVNYLPG QGFAGFRNPPSNRIFACWGKPA
						WTACCNSLRARR
13488 4	43856	В	13568	350	742	
13489 4	43857	A	13569	248	599	HRDSYKRMAYGLVISVVSLSR KLRKPVL*PDKEGNEIWVDMY TVKPSGWTVRTFDKPRKRFIAF FIAGILFRAIKNHFLPRETLQCLF YILTGFRRGQSEYFSIFSNMDLA DTVMFL
13490 4	43858	В	13570	1	2148	
13491 4	43859	A	13571	3064	5069	RSDPDENCC*TGH*STLFNWC\ WRNALSIWEPVCNEIFYRLIKPR WEIRWGKRAPVIPKHTLNTQPV EDTSLSTPAAPMVDSLIARVGV MARGNAITLPVCGRDVKFTLE VLRGDSVEKTSRVWSGNERDQ ELLTEDALDDLIPSFLLTGQQTP AFGRRVSGVIEIADGSRRRKAA ALTESDYRVLVGELDDEQMAA LSRLGNDYRPTSAYERGQRYAS RLQNEFAGNISALADAENISRKI ITRCINTAKLPKSVVALFSHPRQ APCMKSNNALIVILGTVTLDAV GIGLVMPDLALKGLRVLLVEG NDPQGTASMYHGWVPDLHIHA EDTLLPFYLGEKDDVTYAIKPT CWPGLDIIPSCLALHRIETELMG KFDEGKLPTDPHLMLRLAIETV AHDYDVIVIDSAPNLGIGTINVV CAAHALIVPTPAELFDYTSALQ FFDMLRDLLKNVDLKGFEPDV RILLTKYSNSNGSQSPWIEEQIS DAWGSMVLKNVHAIVTGDAV GMDDIPQEARQYRHNQAYAYS IQGDGAEDDDERIVRFHTRVTV DSDTLASDAARLTCRHGLGNQ RNGGVLIEDKFECFFSLTAVFA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13492	43860	A	13572	1408	5578	CSHLLIIQLANQNTIITFGKCSSF
13492	43800		13372	1400	3370	TTATPIGNFYDTRYSSTERRCLL
						TSQ*KRRDEIIQCVLSKQL\RAR
						SLPDPTRKVFSTLSPRSTFRVNF
				[TSRPHTGKVWAFREPFLLPPHL
						RKPDTDTGYLYKRMAYGSVIS/
						ICFTKPKLRKPVL*PDKEGNEIW
						VDMYTVKPSGWTVRTFDKPRK
						RFIAFFIAGILFRAIKNHFLPRET
						LQCLPYILTGFRRGQSEYFSIFS
						NMDLADTVMFL
13493	43861	A	13573	2678	2884	SAPLQRNTARRFLLFLKACSNR
10 170						FCYR\WRHMTAIPSYNRSKLREI
						KGGLYHMGIFRRFLASKETPTF
						CY
13494	43862	В	13574	1 .	7903	
13495	43863	Α	13575	1	880	
13496	43864	Α	13576	1	996	
13497	43865	Α	13577	763	778	NRCEPPV*GTRFSQQLLLALKA
						TLLIEELYALTVHSWRPKW
13498	43866	Α	13578	2	254	
13499	43867	Α	13579	1	2358	
13500	43868	Α	13580	206	382	
13501	43869	С	13581	1	1137	
13502	43870	Α	13582	79	906	
13503	43871	Α	13583	2	383	SPPEVVRLGYHPPRHGHQ\HLF
]				ASCRKGDVGRVRP*TRPDPGSD
						LLPAQVPAGAARRGGTMPACV
İ		}				GTRSWYSTFWPMEPAARPTPS
						MVSAASMGH*VTPSAGLYAITS
						RSRLPAGGGITMTTSCSGF
13504	43872	A	13584	1	1092	
13505	43873	Α	13585	173	262	
13506	43874	Α	13586	898	1391	QLGILRVNLLGIDLRVTGHQAA
						PPLHLLDLLQMH/SSPPCRPQ*A
						TSSPPPSLRCRAPHR*SLAAP*A
						PPPRPSLSAGSPHRKGPPGPAPT
	ŀ					GSWSSGNQAGLPCCLGWASVF
						LCPSHLPPPHLPQSSASLAFHRV
		ŀ				KHLIHLHNSVIDKMKYQLKFLC
		ļ	12.50	<u> </u>	10.00	LLGEYLSV
13507	43875	A	13587	1212	960	DOELING HOLD TO THE STATE OF TH
13508	43876	Α	13588	1318	1369	RDFHPK*HTIGIQYIWQQLEINQ
						CDTSLSMVLP\R*RDFHPKSHLL
12500	42.077		12500	212	205	PGAAQIQ
13509	43877	A	13589	212	385	
13510	43878	A	13590	377 69	505	CSI DSI A OMI DEVDDD SI CI SOS
13511	43879	Α	13591	03	193	CSLRSLAQMLPFYPRRSLCLSCS L**PHCCLWLAALRSESK
13512	43880	A	13592	1	230	L FRECEWLAALKSESK
13512	43881	A	13593	2	341	
13313	143001	I ^A	13333		1 1 1	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13514	43882	Ā	13594	1	2421	
13515	43883	Α	13595	1	623	
13516	43884	Α	13596	1	906	
13517	43885	A	13597	1	568	
13518	43886	A	13598	107	199	
13519	43887	A	13599	42	328	GGASGVCSF*CSE/LSSFWWVR
	1,500,	1	13377			GLAGSGVKLQTFATQEPSWLH
						PVDPVPGLQVELPASPAPCART
						PQPLGGRWDSAPWSRGRRSSG
						AHGIDVALLP
13520	43888	A	13600	5	124	IWILWFFVSLT*RSRGPSQ*VLQ
13320	43000	\\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	13000]	124	LIKAPHVDLLVVTFGC
13521	43889	A	13601	827	973	IHQSALCI*LKFYTSLTHPVISLV
13321	43009	A	13001	02/	973	VETLVWW*YLKM*KKLWSWI
						RGRG
12522	12000	<u> </u>	12602	2	502	RORO
13522	43890	Α	13602	2	503	
13523	43891	Α	13603	56	792	
13524	43892	A	13604	1	411	
13525	43893	A	13605	1	1554	
13526	43894	A_	13606	1	363	***
13527	43895	Α	13607	1	966	
13528	43896	A	13608	932	1189	
13529	43897	Α	13609	146	323	
13530	43898	Α	13610	1	1827	
13531	43899	Α	13611	2	412	
13532	43900	Α	13612	1	2238	
13533	43901	Α	13613	2	411	GATDCVCYYTVGFNDTKTSAL
		1				HMVVGDSLAMDVSSVHHNSTL
						LRYSVTLLGYGFYGDIIKDSDK
		1				KRRLGLARYDFLCIKTFFSHH*
	ł				ł	YEGTVTFLPAQHTVGSPRDRKP
						CRAGCFVCRQSKQQLEEEQKT
	J					ALYGLEL
13534	43902	Α	13614	3	2185	
13535	43903	Α	13615	206	1568	
13536	43904	Α	13616	165	417	TWDYIRPNE*YM*MAKKKMP
						MD\VWCNGKNWRPAG\EFVDD
						G\TETHFSIGNHDCYIKAVSSGK
						RKEGIIHTLIVDNREIPEIAS
13537	43905	Α	13617	463	837	
13538	43906	A	13618	1	765	
13539	43907	Α	13619	3	510	AGVHKIEFEHGTTSGKRVVYV
						DGKEEIRKEWMFKLVGKETF\Y
					1	VGAAKTKATINIDAISGFAYEY
						TLEINGKSLKKYMEDRSKTTNT
						WVLHMDGENFRIVLEKDAMD
						VWCH\GKKIRRPAGEFVDDGTE
						TPLQYPGPMTCYIKAV\SSG\KR
						KEGIIHTLIVDNREIPEIAS
13540	43908	A	13620	342	585	
	43909	A	13621	1	1119	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
13542	43910	В	13622	48	1104	
13543	43911	A	13623	656	916	GWRKYLQGAADMGIAEGGGS VWSSWASSSLISVPVR*LCESTQ RSAPGWPPLTGGSCSESVALWL CVNDFEFQMRLMPSRDSEPAC
13544	43912	Α	13624	179	682	
13545	43913	A	13625	1	7511	
13546	43914	Α	13626	56	8669	
13547	43915	Α	13627	42	8716	
13548	43916	Α	13628	1	2457	
13549	43917	Α	13629	3	571	
13550	43918	Α	13630	1	867	
13551	43919	Α	13631	1	2238	
13552	43920	Α	13632	171	396	
13553	43921	Α	13633	1	3940	
13554	43922	Α	13634	554	619	
13555	43923	Α	13635	1	1359	
13556	43924	С	13636	208	359	
13557	43925	A	13637	2	513	PSHDHSKLGTRASGKVILHGEH AAVHGKVALAVSLNLRTFLRL QPHSNGKGDLSLPNIGIKRAWD VARLQSLDTSFL/RWTKEDLELI NKWAFQGERMIHGNPSGVDNA VSTWGGALRYHQGKISSLKRSP ALQILLTSTRG\PRNIRALVGGG RNRL\LEVPDMGGPLLLSW
13558	43926	A	13638	3	1331	RVDDFVGELLRLRRGGAADVE AAAAAGFPGAMLSQVLLVSAP GKVILHGEHAVVHGKVALAVS LNLRTFLRLQPHSNGKVDLNLP NIGIMRAWDVARLHSLDTSFLG ECKEEKPEQGDVTTPTSEQVEK LKEVAGLPDDCAVTERLAVLA FLYLYLSICRKQRALPSL\DIVV WSELPPGAG\LGSSAAYSVCLA AALLTVCEEIPNPLKDGD\CVN R\WTKED\LELINK\WAF\QGER MIHGNP\SGV\DNAVQHLGEGA LRLPSRGRFSFL*RGSPVFQI/LG LTNTKVPSAITRALVAGVRNRL LKFPEIVAPLLTSIDAISLECERV LGEMGEAPAPEQYLVLEELIDM NQHHLNALGVGHASLDQLCQV TRARGLHSKLTGAGGGGCGITL LKPGLEQPEVEATKQALTSCGF DCLETSIGAPGVSIHSATSLDSR
13559	43927	Α	13639	1	950	
13560	43928	Α	13640	1	1653	
13561	43929	Α	13641	1	2601	
13562	43930	Α	13642	2	513	
13563	43931	Α	13643	47	367	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13564	43932	A	13644	1	633	
13565	43933	В	13645	1	1110	
13566	43934	Α	13646	172	1095	
13567	43935	A	13647	1	855	
13568	43936	Α	13648	3	440	
13569	43937	Α	13649	2	230	
13570	43938	A	13650	2	606	AEFDLCCSPCRRRLLGREEAGE EPTSPVTQYLQPRSPEECKMFA CAKLA\CTP\SLIRAGSRVAYRPI SASVLSRPEASRTGEGSTVFNG A\QNGVSQ\LIPKGSFKTSAISKK TLDTA\AKFIG\AGAGNS*GVG WFLVAGIG\TVFWASLIHWVM ARKPFRWKQQL\FSY\AILGICL VLKAMGSPFVLMVAFLILFAHV TEITA
13571	43939	A	13651	200	2320	1
13572	43940	A	13652	224	640	
13573	43941	A	13653	1	357	
13574	43942	A	13654	43	183	
13575	43943	A	13655	3	478	SSAGREPDPSDLPRRLCFTHRLP AARRWVQLCVHASPEPGGQGV
						CPGRSERMVIRVFIASSSGFVAI KKKQQDVVRFLEANKIEFEEVD ITMSEEQRQWMYKNVPPDKKP TQGNPLPPQIF\NGDRYCGDYD RLFESKESNTVFSFLGLKPRLAS KAEP
13576	43944	A	13656	2	428	CRDGKDVVSLVRATVWALSKR KLQPTRA/ALTPTPSAVNKIKQF LKDKPELVDVKVGVRTRGCNG LSYTVEYTKTKGNSDEVIQDEA RVFIEKKAQLTPLGTEMDYVED KLSSE/FMFNNPT*TCGCGAPGK LVETLGLTEEIM
13577	43945	Α	13657	1	1377	
13578	43946	Α	13658	1	1215	
13579	43947	A	13659	1	452	REAEAADLRSEAPGRRDGEDV GFLSSGQLSGLCSKRKLQPTRA ALTLTPSAVNKIKQLLKE*P*AC R\LKVGVRTRGCNGLSYTLEYT KTKGDSDEEVIQDGVRVFIEK\K AQLTLL*TEMDYVEDKLSKGV GVH\SPNIKGTCGCGESFNI
13580	43948	Α	13660	3	763	
13581	43949	Α	13661	1	306	
13582	43950	A	13662	738	1022	GQHHPETKAWQRHNKKREF*T NILDEH*CKNPH*NTGKPNPAT HQKAYPP*SSGLHPWDARLVQ HTKINKCNPAYKQNQRQKPHD YLNRCRKGL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13583	43951	A	13663	2	516	QKIDLPEYQGEPDEISIQKCQEA
						VRQVQGPVLVEDTCLCFNALG
						GLPGPYIK\WFLEKLKP\EGLHQ
1						FLAGVEGKS\AYALCTFALSTG
						DPSQPVRLFRGRTSGRI\VAPRA
						CQDFGWD\PCFQPDGYEQTYAE
		1				MPKAEKNAV\SHRFRA\LLE\LQ
		L.				EYFGKFGQLDFCKLEAGPFR
13584	43952	Α	13664	7	227	
13585	43953	Α	13665	1	415	
13586	43954	Α_	13666	234	510	
13587	43955	A	13667	2	402	
13588	43956	В	13668	530	1544	
13589	43957	A	13669	1	603	
13590	43958	В	13670	1	1527	
13591	43959	A	13671	1	1179 413	
13592	43960	Α		2	374	KHISPALKALELPFERNLII/PLQ
13593	43961	A	13673	100	374	LLKVRILKEGEML/DI*HWFL**
		1				GMGTVQKGMPHKCYHGKTGR
						VYNVTQHAVGIVVNKQVK*VV
13594	43962	A	13674	1	1035	VINVIQUAVGIVVINQVR
13595	43963	A	13674	2	564	GPFGRNRHLPVIRQNDGPQRGK
13393	43903	^	13073	2	304	RRGTPKYVF*/RPFRK\HGGGPL
		1				GPPYM\RI\YRKGDIVD\IKGMG\
İ						TVQNGMPH\KCYHGKTGRVYN
						VTQHAV\GIVVNK\QVKGKILA
		1		:		K\RINVRIEHI\KHSKSPR*ASLK/
						RVLKENDQEKERSPKKKGTWG
		1				SNLKRHPAPPQKKHTL*RTNGK
		İ				EPEL\LEPIPYEFHGHN
13596	43964	В	13676	68	391	
13597	43965	Ā	13677	1	404	PTRTWTRGRIPRLSAPPSRGAR
						GTMADPRVRQIKIKTGV\VKRL
						V\KEKVLYEKEPKQ\QEEKIEKM
						RAEDGENYDIKKQAEIL\QESR
						MMIPDCQRRLEAAYLDLQRILE
						NEKDLE\EAEEYKEARLVLDSV
						KLEA
13598	43966	Ā	13678	1	436	LLLACRSMDCYLAIVHATRTLT
						QQRHLVKFICLGLRNLFLLLSL
		1				RILLFRRTFYPSNVSPVCYEDM
						GNNTANWWMLLRILPQSFGFIV
						PL/LIMLFCYRFTLHTLFKAHMG
					1	QKHWTMWVIFADVLIILLCWL
						PYNLVLLADTIMGT
13599	43967	Α	13679	1	705	
13600	43968	Α	13680	292	1489	
13601	43969	Α	13681	434	1205	
13602	43970	Α	13682	1	390	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13603	43971	Α	13683	21	479	TYPEAWKLYYRSR*SRDLRKY
						HAHSVSP*QDQRYTS*KILKCD
		1				EHIQKLGSCITDPGNRETSGNT
		İ				MHTVFHRDKTKDTHPESCCSSE
						KGGQPLPWFEHRKNVPQFAEP
						TETLFGPDSGKGAKSLVELLDE
						SECTSDEEIFISQDEIQSFMANN
		İ				QSFYSNREQYRQHLKEKFNKY
					•	CRLNDHKRPICSGWVDNGWEA
13604	43972 .	В	13684	32	438	
13605	43973	Α	13685	533	1226	
13606	43974	Α	13686	1	1566	
13607	43975	Α	13687	215	453	
13608	43976	Α	13688	1	553	RRPARAAVIKQPPAPGASQHAA
						TPTQTPLCTPARPLPPMAHVQQ
						LEGRWRLVDS\KGF**NTMKE\L
						GVT\IALRKMGRNAQAQICIIT\\$
						DG*NPSPLKTESTLKT\TQFSCT/
						LWGEKFEETTA\DGQKKLTNCL
						STFPDG\AL\VQHQEWDGERKS
						T\ITKKN*KDGEISWLECVMNN\
						VTCTPDSMKK
13609	43977	С	13689	11	106	
13610	43978	Α	13690	3	155	LCELQKAIDLFTDAIKLNPRLAI
						LYAKRASVFVKLQKPNAAIRD
						CDRAIEINPDSAQPYKWRGKAH
						R*ILAWPFCMPRGPVSSSNYRS
			<u> </u>			QMLPSETVTEPLK
13611	43979	Α	13691	1	1732	
13612	43980	В	13692	95	1715	
13613	43981	Α	13693	1	446	
13614	43982	A	13694	1	786	
13615	43983	A	13695	88	828	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
13616	43984	A	13696	157	1515	RREKMAELKYISGFGNECSSED
						PRCPGSLPEGQNNPQVCPYNLY
						AEQLSGSAFTCPRSTNKRSWLY
						RILPSVSHKPFESIDEGHVTHNW
		ŀ				DEVDPDPNQLRWKPFEIPKASQ
		ŀ				KKVDFVSGLHTLCGAGDIKSN
		l				NGLAIHIFLCNTSMENRCFYNS
						DGDFLIVPQKGNLLIYTEFGKM
						LVQPNEICVIQRGMRFSIDVFEE
		İ				TRGYILEVYGVHFELPDLGPIG
		ļ				ANGLANPRDFLIPIAWYEDRQV
	1					PGGYTVINKYQGKLFAAKQDV
						SPFNVVAWHGNYTPYKYNLK\
						NFMVINSVAFDHADPSIFTVLT
	1					AKSVRPGVAIADFVIFPPRWGV
						ADKTSGPPYYHR\NCMKRFME
						LIRGQFEQKQGWFLPGGRG/SL
						HSTMTPHGP\DADCFEKASKVK
						LALKRIADGTMA\FMF*SSFNLA
						GPKWGLQAP\RVLDKNYPKCW
						EPLKSPF\TPNSRNPAEPN
13617	43985	Α	13697	1	156	
13618	43986	Α	13698	8	306	
13619	43987	С	13699	1	1143	
13620	43988	Α	13700	1	1551	MRKDSCASSMHQQVSRSKKRA
ŀ		1				GQKTPPEDQEGGQRALRSSHIR
						LGQFLLIEDCKTPSPSSLGADAI
						AKQRKTSVSAAASVSATIPIRR
1						VQGPTVVGSWARGVSAAASGP
1						RGTGPKGKARSEKGCSLSHGPQ
1						TNKPLVVQKGQKMEQANHPV
						GLVISVVYKDILKKIVQRETSHP
						LIHVRYAEAITGRRTAPEDKGS
						LGRDMLAKAGAIIYMNMGNKL
						PIWCHLLEEGIYLEVWALEGQF
1						GRAKNACPVQIRLKDPTTFPYQ
		ŀ				RQYPLRPEAHKGLQDIVKHVK
ŀ						AQGLVKKCSSPCNTPILGVQKP
		ŀ				NGQWSLVQDLRLISEAVIPLYP
ŀ						VVPNPYTLLSQIPEEAEWFPVL
ŀ					}	DLKDAFFCIPLHYDSHDSQFLF
						AFEDPTDHTSQLIWTVLPQGFR
						DSPHLFGQALAQDLGHFSSPGT
		1				LVLQYVDDLLLATSSEASCQQA
						TLDLLNFLANQGYKASRSKAQ
	1]			LCLQQVKYLGLILARGTRTLGK
						ERIQPILAYPHPKTLKQLWGFL
	100	<u> </u>	1222	1.00	600	QIT/GFCQLWIPR*SKI
13621	43989	A	13701	188	688	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	:	
13622	43990	Α	13702	636	2628	SNDRTEDDCGKHPFMSSPP\TEP
					:	WVCLIIEGQEIDFLLDTGTTFSV
ļ						LIPCLGRLSSRSVTIQGILGQPVT
						RYFSHLLSCNWETLLFSHAFLV
						MPESPTPLLGRDILAKAGAIISM
]		KTGNKLPICCPLLEGINPEVWA
						LEGQFGRAKNAHPLQIRLKDPI
;						SFPYQRQYPLRPEAHKGLQDIV
						KHLKAQDSVRKCSSPCNTPILG
		1		1		VQKLNSQWRLVQDLRLINEAVI
						PLYPVVRNPYTLLSQVPEEAEW
10.600	10001		12502	202	(10	FTVLDLKD
13623 13624	43991	В	13703	302	619 5172	
13624	43992 43993	A A	13704 13705	1	2091	
13625	43994	A	13706	1	1754	GPRGTGPKGKARSEKGCSLSHG
13020	43774	l'a	13700	'	1754	PQTNKPLVVQKGQKMEQANHP
						VGLVISVVYKDILKKIVQRETS
						HPLIHVRYAEAITGRRTAPEDK
						GSLGRDMLAKAGAIIYMNMGN
						KLPIWCHLLEEGIYLEVWALEG
						QFGRAKNACPVQIRLKDPTTFP
						YQRQYPLRPEAHKGLQDIVKH
						VKAQGLVKKCSSPCNTPILGVQ
						KPNGQWSLVQDLRLISEAVIPL
		l				YPVVPNPYTLLSQIPEEAEWFP
					<u> </u>	VLDLKDAFFCIPLHYDSHDSQF
						LFAFEDPTDHTSQLIWTVLPQG
		l			1	FRDSPHLFGQALAQDLGHFSSP
						GTLVLQYVDDLLLATSSEASCQ
						QATLDLLNFLANQGYKASRSK
						AQLCLQQVKYLGLILARGTRTL
						GKERIQPILAYPHPKTLKQLWG
						FLEITGFC*LWIPGYSKIARPLYT
			:			LIKETQRANTHLVEWESEAETA
			:			FKTLKQALVQAPGLSLPTGQNF
						SLYVTERAGIALGVLTQTRGTT
						PQPVAHLSKETDVVAKGWPHC
						LRVVAAVAVLVSEAIKIIQGKD
						LIVWTTHEVNGILG/AKGSLWL
						SDKRLFRYQALCLEGLVLQIRT
		_				SCGSHLAVTRLWALYF
13627	43995	Α	13707	2	887	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		:		sequence		
13628	43996	I A	13708	3	1750	GWAVPRPGPRGLTVATEAAAA
13020	43990	^	13708	3	1750	AAAAAAAAISSRTRAPQAPEGR
						NRRLEKMADDIDIEAMLEAPY
						l l
						KKDENKLSSANGHEERSKKRK
						KSKSRSRSHERKRSKSKERKRS
						RDRERKKSKSRERKRSRSKERR
						RSRSRSRDRRFRGRYRSPYSGP
1						KFNSAIRGKIGLPHSIKLSRRRS
						RSKSPFRKDKSPVREPIDNLTPE
						ERDARTVFCMQLAARIRPRDLE
						EFFSTVGKVRDVRMISDRNSRR
						SKGIAYVEFVDVSSVPLAIGLTG
						QRVLGVPIIVQASQAEK\NRAA
						AMANNLQKGSAGPMRLYVGSS
						HFNITEDMLRGIFEPFGRIESIQL
						MMDSETGRSKGYGFITFSDSEC
		l				AKKALEQLNGFELAGRPMKVG
		l				HVTERTDASSASSFLDSDELER
ĺ						TG\IDLGTTG\RLQLM\ARLAEG
						TGLQIPPAAQQALQMSGSLAF\
						GAVGRNLFCYRFAKQDFSHQT\
						ETSALSAAAS\VQPLATQCFQLS
						NMFNPQTEKEVGWDTEIKDDV
						IEECNKHGGVIHIYVDKNSAQG
		l				NVYVKCPSIAEAIAAVNALHGR
						WFAGKMITAAYVPLPTYHNLF
1						PDSMTATQLLVPSRR
13629	43997	A	13709	141	343	LHEGLLP*RGHALLSGALHG*C
1302)	13777	l'`	1370)	' ' '		VGSCLFSTPRCLTSSTACGYPSS
		1				CTPGSCTAWCPSASRRLSCWT
13630	43998	Α	13710	5	426	
13631	43999	Α	13711	1	1188	
13632	44000	Α	13712	2	264	NKQPIWIPSRHLKPYHEPDAKE
1		1				EIPGGS*GPTSCSHVETDAEEDP
						NCHEQHLSNTATHLGTDQEAVI
1						DGRRKPEESRTTSHICRCRS
13633	44001	Α	13713	1	700	
13634	44002	Α	13714	3	483	
13635	44003	Α	13715	2	361	RRFRAGAGYVVESTGVFTTME
						KAGAHLQGGAKRVIISAPSADT
						PMFVMGVNHEKYDNSLKIISNA
		1				SCTTNCLA/PPASTGAAKAVGK
						VIPELNGKLTGMAFRVPTANVS
						VVDLTCRLEKPA
13636	44004	A	13716	2	249	SADAPMFDMGVNHEKYDNSL
.5556		`				NII/SVMKAGPVEKRPAWHPMD
						TLP*LAPRSLFLCSNASCTTNCL
						EPLAKVIHDNFGIVEGLMV
L	<u> </u>	<u> </u>	L	L	L	EL EVICA HIDIA OLA FORMA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	ſ	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13637	44005	A	13717	1	1079	GTRRQSAASSFASPAEPHRSDT
į						MGKVKVGVNGFGRIGRLVTRA
						AFNSGKVDIVAINDPFIDLNYM
						VYMFQYDSTHGKFHGTVKAEN
						GKLVINGNPITIFQER\YPSKIKW
						GDAGAEYFVESTGVFTTMEKA
						GAHLQGGAKRVIISAPSADAPM
	ļ					FVMGVNHEKYDNSLKIISNA\S
		ļ				CTTNCLT\PLAKVIHDNFGIVEG
						LMTTVHAITATQKTVDGPSGKL
						WRDGRGALQNIIPASTGAAKA
						VGKVIPEL\NGKLTGMAFRVPT
	ŀ					ANVSVVDLTCRLEKPAKYDDI
	ŀ					KKVVKQA\SEGPLKGILGYTEH
	ŀ	1				QVVSSDFNSDTHSSTFDAGAGI
						ALNDHFVKLISWYDNEFGYSN
				1		RVVDLMAHMASKE
13638	44006	A	13718	3	483	
13639	44007	Α	13719	2	361	RRFRAGAGYVVESTGVFTTME
						KAGAHLQGGAKRVIISAPSADT
						PMFVMGVNHEKYDNSLKIISNA
		ŀ				SCTTNCLA/PPASTGAAKAVGK
		ľ				VIPELNGKLTGMAFRVPTANVS
		į				VVDLTCRLEKPA
13640	44008	Α	13720	2	249	SADAPMFDMGVNHEKYDNSL
						NII/SVMKAGPVEKRPAWHPMD
		ļ				TLP*LAPRSLFLCSNASCTTNCL
						EPLAKVIHDNFGIVEGLMV
13641	44009	Α	13721	1	1079	GTRRQSAASSFASPAEPHRSDT
		ŀ				MGKVKVGVNGFGRIGRLVTRA
						AFNSGKVDIVAINDPFIDLNYM
	,					VYMFQYDSTHGKFHGTVKAEN
						GKLVINGNPITIFQER\YPSKIKW
		ŀ				GDAGAEYFVESTGVFTTMEKA
					:	GAHLQGGAKRVIISAPSADAPM
						FVMGVNHEKYDNSLKIISNA\S
						CTTNCLT\PLAKVIHDNFGIVEG
						LMTTVHAITATQKTVDGPSGKL
		1				WRDGRGALQNIIPASTGAAKA
						VGKVIPEL\NGKLTGMAFRVPT
						ANVSVVDLTCRLEKPAKYDDI
1		1				KKVVKQA\SEGPLKGILGYTEH
						QVVSSDFNSDTHSSTFDAGAGI
						ALNDHFVKLISWYDNEFGYSN
						RVVDLMAHMASKE
13642	44010	С	13722	40	243	
13643	44011	Α	13723	2	2972	
13644	44012	Α	13724	1	602	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13645	44013	A	13725	1	325	MGRNQSRKAENTKNESASSPP
						KDRNSSPVREQSWTENEFDELT
	III					EVGFRKLVITNFSELKEHVLTD
1		ŀ				HKEAKNLEKS\LEVVAREISQE
		e e	ļ			KEIKVFN*EKRKSNCPCLQMT
13646	44014	A	13726	1	981	
13647	44015	Α	13727	1	739	MGRNQSRKAENSKNQSTSSPPK
						DCSSSPAREQNWTKNEFDELTE
						VGFRRSLIANFSKLKECVLTHR
						KEAKNLEKSDRENGTKLENTL
						QDIIQENFPNLARQANIQIQEIQ
			!			RIPQRYSLRRATPRHIIVRFTKV
		ŧ				EMKEKMSRAAREKGRVTHKG
			[KPIRLTADLSAENLQARREWGP
	ľ					VFNILK\NFQPRIPYPAKLSFISG
						GEIKSFTDKQMLRDFFTTRPAL
						QELLKEALNMERNNQYKPLQK
						HAKW
13648	44016	Α	13728	1	1277	MKEQNGGGGREEDHKDKGIRQ
						LPSLLPSAGEAKSDKPEHHNRY
					,	RKEIASFEDEKRAMGVAKMAE
}						QEQLQSAAPSMTNAEVGTAHG
						GRAEAGRGVDSPGKRKVLGDF
						PFLAKGSRDRLPGKAGHSLPKY
						YAFTKVLATGRQERNSINISKK
						VIYTKTTSTGHQHQRPKIDKTT
		ŀ				KMGRNQSRKAENSKNESTSSPP
						KGHSSSPATEQSGTENDFDELT
		:			;	EVGFRRSVTTNFYELKEDVQTH
						RKEAKNLEKRLDKWLTRINSV
						EKTLSDMMELKTMARELRDTC
i						SSFKSRFNQVEERVSVIEDPIY\E
ŀ						LSREDKVRDKKVRNKQSLQEI
						WDYVKRPNLRVIGVPESDGEN
		1				GTKLENMLQDIIQENFPNLARQ
						ANIQIQEIQRTPQRYSSRRATPR
						HIIVRFTKVEMKQKVLRAAREK
10540	14015	 	12722		1504	GPVTHKGNSGSLS
13649	44017	A	13729	1	1584	CHI DAMIDA NITO CODNICTIVIZZO
13650	44018	A	13730	493	575	SHLPMNPANTRSQRNPTKKKQ
i						KKNVC/DNIRELEQQMEDAYR
						GTKRKMLPSSSSRMRSDGFDEE
						SQRYYWRPKNEISGTLEDDFLK
						AKSWNRKFYDYEANMPDRWG\
						QVAIKSYTLKNLKQTVVISKILP
						TGRKHLPR*SHLPMNPANTRSQ
12651	44010	 	12721	1	655	RNPTKKKQKKNVCG
13651	44019	A	13731	1	655	
13652	44020	A	13732	2	1664	ļ
13653	44021	A	13733	1	2364	
13654	44022	A	13734	1	224	

1	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13655	44023	A	13735	1033	7511	
13656	44024	Α	13736	56	8669	
13657	44025	Α	13737	42	8716	
13658	44026	Α	13738	1	5046	
13659	44027	Α	13739	1	2298	
13660	44028	A	13740	1	2362	
13661	44029	Α	13741	1	2046	
13662	44030	Α	13742	1	1503	
13663	44031	Α	13743	I	813	
13664	44032	A	13744	185	735	
13665	44033	В	13745	317	910	
13666	44034	Α	13746	1	867	
13667	44035	В	13747	1	2325	
13668	44036	Α	13748	1	430	
13669	44037	Α	13749	1	867	
13670	44038	A	13750	35	518	CLSRRASPVYLASMSGRGKTG GKARAKAKSRSSRAGLQFPVG RVHRLLRKGHYAERVGAGAPV \YLAA\VLEVAQPLRVLELAGN AA\RSN\KKTRIIPRHLQLAIRND EEL*KLPGLP*RIAQGSR/VLPNI QA\VLLPK\KTSATVG\PKAPSG GKKATQASQEY
13671	44039	Α	13751	1	937	0
13672	44040	A	13752	1	1096	
13673	44041	A	13753	85	1512	
13674	44042	A	13754	3	2377	
13675	44043	Α	13755	1	1134	
13676	44044	A	13756	49	408	
13677	44045	Α	13757	47	2603	
13678	44046	A	13758	1	378	
13679	44047	Α	13759	1	696	
13680	44048	A	13760	3	401	
13681	44049		13761	882	1106	KLHPYTTRQYCHFRI*TDVMAK RSRNAVKGLSALVWCALSDFA AGFTPDDGCTRATFFTGGWRK RDCRGAVAGG
13682	44050	Α	13762	688	1100	
13683	44051	Α	13763	476	552	
13684	44052	В	13764	178	1107	
13685	44053	Α	13765	1	198	
13686	44054	В	13766	94	625	
13687	44055	Α	13767	1	732	
13688	44056	Α	13768	1	4443	
13689	44057	Α	13769	216	575	PGAGDRYVGKPVSAGVTAVSC GVWSSMPDATLIASYQAYTSH VGRIRRSRRIR/TNTARADYPAL TALPFNNFRTHNRFGFNAASVL ASIGIGSSLICAARISAHSGAEH KPREPKAANIK

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	1100	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13690	44058	A	13770	1018	1489	KLPGQRRFVQTPSHVLHTTLSY
						PTAPAQSSYPAPLSPSQDRAPTL
						AHQTT*FSASYTANAQWQRVA
						ADHLKSVRETYAHVRQFSPAL
						GNAPPDLQPLS*ASCVPKSGAR
						YSSVIRQNRRDFFRATDHMYVT
						VPTNRPKMWNPVNTQMNSFRA
						DTCER
13691	44059	A	13771	1013	1755	GSPGLRPGGAAGKALAPSGCT
						GNSHRAPM*PGAQAQTRPEKW
						A*QRP/HVSTGARGARVRALAA
						VAAAEAQRAGAAVRAAALHT
						RAAVSAGAARAVVQVVLAAR
						TREAGAAAAAQGVAQVQAES
						AIAAWLQAAPVHPLLTVGALE
						TGWAVADVGRVRVCTPDTQA
						AVEAGSICTCHPAHLTPQPVEP
						TRAGAFKGPGCLTTAASIGTRV
						PVTGPRPRDKASGGSRGRAPGA
13692	44060	A	13772	1	1236	EAAAGGDRDRAEGSSPAL
13693	44061	A	13773	1	456	
13694	44062	A	13774	11	219	
13695	44063	Α	13775	1707	1853	
13696	44064	A	13776	1	615	QLATFFFGGGLPGGSAPWDPPG
		1				EACLWVLVRLPGRTVWVIQFV
						QLRSFFAPPFAHNGRASSGRVS
						VHYGIAQTAEWAAEICGVGAR
				İ		KIRELAAIFHQNTIMHFANGG/N
						PTRRSAVLSSMQGSLPGGC\DA
						VDIHPVARIVEALENPGGAYQH
	İ					NGMNRHFPDIRFIWWAGGANF
						THHQDTNRLIRA WQNDSYFRK
						GRIKAKMGPKLF
13697	44065	Α	13777	407	661	ANWIKWRWHRAMHSSSSMWQ
		ŧ				TANSLASFISAPVTSSSACRSTL
				ŀ		PATRYWCI*WRSSAIWKWVILS
10.500	111055	<u> </u>				GPVATRICTATIWIQLICN
13698 13699	44066 44067	A	13778 13779	3	2448 1612	
13700	44068	A	13780	1	1626	
13701	44069	A	13781	3	118	
13701	44070	A	13782	1175	1396	
13702	44071	A	13782	197	933	
13704	44072	A	13784	2	64	IILLNSAPAWM*TFKLGPGVDV NYMLHTQNK
13705	44073	Α	13785	1	855	
13706	44074	В	13786	1	1605	
13707	44075	A	13787	1	588	
13708	44076	Α	13788	1	2769	
13709	44077	Α	13789	813	1341	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13710	44078	В	13790	1	999	
13711	44079	Α	13791	1	281	
13712	44080	Α	13792	1	1332	
13713	44081	A	13793		1384	MPITLRRSVEKEQLIEIANTIMPF GKYKGRRLIDLPEEYLLWFARK DEFPAEARYTHIDLHVPAPLLV MFTLDTRQSACDRNPHLPIHVC SSWTSPKCAPRLWTRALLPRTR TPPYATGTTIGSRASWFVTFPD ASFYRKAPRQEFHTSLQGRSL YVRQRSTANPTEITRRTLIHKM VEVNACLKQLDNKDIADYEHN QLMRRLRQLIAQSWHTDEIRKL
12714	44000		12704		2004	RPSPEAIMTEQEKTSAVVEETR EAVDTTSQPVAT MTQQITLIKDKILSDNYFTLHNI
13714	44082	A	13794		2004	TYDLTRKDGEVIRHKREVYDR GNGATILLYNTKKKTVVLIRQF RVATWVNGNESGQLIESCAGL LDNDEPEVCIRKEAIEETGYEV GEVRKLFELYMSPGGVTELIHF FIAEYSDNQRANARGGVEDEDI EVLELVQCGGQIGVPDTVLAVF ATGIGFLAMSVTKPRINAQPYP MPWRNLAQLVEHINRTRIHRNL QFTDAFKRRLIDNICRKNNIIGT HTGQSKFSGQTQQIFGEIPQVVP ASELAAAMKQIKELQRLLGKK TMENELLKEAVEYGRAKKWIA HAPLLPGDGERTDDWMDGRRS RHTDDTDVLLRIHHVIGELPTY GYRRVWALLRRQAELDGMPAI NAKRVYRIMRQNALLLERKPA VPPSKRAHTGRVAVKESNQRW CSDGFEFCCDNGERLRVTFALD CCDRLSSRGRNAAAKRQRDDP LRWKFRLPSGGTKLLMPLDQQ APARFAPFVTCTLTILCGTGHV VYTILPIIYDVAIKNNIRPERPM AASSIGAQMGIIASPVSVAVVSL VAMLGNVTFDGRHLEFLDLLAI TIPSTLIGILAIGIFSWFRGKDLD KDEEFQKFISVPENREYVYGDT ATLLDKKLPKSNWLAMWIFLG AIAVIALLGADSDLRSSF/GGKP

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	•	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				l l l l l l l l l l l l l l l l l l l		
13715	44083	Α	13795	3	918	DKTRKRRIRQ*IPNATSFLYQNS
						PNGLSE*L\PAGTWFAGNAPASS
						CGFRDRYLIVCSHQTPHVPGQI
						TVAAPDARAQAILRIVRQLQRF
		ļ				FHRFEGGHRQHRAKNLLLEHP
						HIVLTKONRRFEADALOPDPA
		Ì				WQQGTLSNGLQWQVLTTPQRP
						SDRVEIRLLVNTGSLAESTQQS
		ŀ				GYSHAIPRIALTQSGGLDAAQA
						RSLWQQGIDPKRPMPPVIVSYD
						TTLFNLSLPNNRNDLLKEALSY
						LANATGKLTITPETINHALQSH
	ŀ	ŀ	1			DMVATLVFRYLKKGWWRYRL
						KGSTLLSHDPAHPLKQPVEAEK
13716	44084	A	13796	1	1368	MCPIEETASSFGGKPLSMVLVIQ
						MFMLLTGALIIILTKTNPASISK
						NEVFRSGMIAIVAVYGIAWMA
						ETMFAPACYGYYILPTYPSDLA
						AIQFDRSGTTHIGRFVINHSFILP
		ŀ				GLIGVSVSCVFGWIFAAIKRDA
		ļ				AAGRAKENVIFHHFPFQSVKAD
ļ		ŀ				IATNPFKRPGGAFIRITQTFRTV
						QTFRLSAYRLDFAGDRLRISTPR
						AKMRTAFKKDHLRQRRRCIRQ
						RAPPARHNLVGAVALPATVAG
						VNFTFSNVPLDSSVLSSLLTDFS
						TAVGSIVMLAVIMGLMLAFDM
						GGPVNKVAYAFMLICVAQGVY
						TVVAIAAVGICIPPLGMGLATLI
						GRKNFSAEERETGKAALVMGC
						VGVTEGAIPFAAADPLR/PSFGG
						KPLSMVLVIQMFMLLTGALIIIL
						TKTNPASISKNEVFRSGMIAIVA
						VYGIAWMAETMFAPACYGYYI
						LPTYPSDLAAIQFDRSGTTHIGR
]				FVINHSFILPGLIGVSVSCVFGWI
						FAAIKRDAAAGRAKENVIFHHF
						PFQSVKADIATNPFKRPGGAFIR
						ITQTFRTVQTFRLSAYRLDFAG
		ļ				DRLRISTPRAKMRTAFKKDHLR
		ł				QRRRCIRQRAPPARHNLVGAV
		ł				ALPATVAGVNFTFSNVPLDSSV
		1		-		LSSLLTDFSTAVGSIVMLAVIM
						GLMLAFDMGGPVNKVAYAFM
						LICVAQGVYTVVAIAAVGICIPP
						LGMGLATLIGRKNFSAEERETG
13717	44085	A	13797	3	2185	
13718	44086	A	13798	1	920	
13/18	144000	l _v	06/51	1	920	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13719	44087	Α	13799	251	479	TQRVDWRSRSSISPGDGLHAFP
						AFGKEGGKMLDRLRERRDVKT
						LQQSGSAAGAEFALAGAHPHT
						GTTA*IQQAINV
13720	44088	A	13800	1	939	
13721	44089	Α	13801	1	386	
13722	44090	Α	13802	3	524	
13723	44091	Α	13803	1	548	
13724	44092	В	13804	67	1144	
13725	44093	Α	13805	811	974	VSRRIKLDVTEEFARAGDNKRC
						ICQ*RHGLAERDIARLLVEIVVI
						WCDIGVIQA
13726	44094	Α	13806	1164	1262	
13727	44095	Α	13807	2	894	
13728	44096	Α	13808	1	1587	MVKEGTFREDLFYRLNVIHLIL
İ				1		PPLRDRREDISLLANHFLQKFSS
						ENQRDIIDIDPMAMSLLTAWSW
						PGNIRELSNVIERAVVMNSGPII
				}		FSEDLPPQIRQPVCNAGEVKTA
	ļ					PVGERNLKEEIKRVEKRIIMEVL
						EQQEGNRTRTALMLGISRRAL
						MYKLQEYDATGFFRDGMTIMV
	İ					GGFMGIGTPSRLVEALLESGVR
						DLTLIANDTAFVDTGIGPLIVNG
						RVRKVIASHIGTNPETGRRMISG
		ĺ				EMDVVLVPQGTLIEQIRCGGAG
						LGGFLNPTGVGTVVEERQTSID
				ł		TSTRKRPFELGRPSRNNDKGIRP
}	•					\DQALDRKNIRE/ALHDSLKRLQ
1				-		TDYLDLYQVHWPLRPFYCFGK
1				-		LGYSWTDSAPAVSQLDTLDAL
				-		AEYQRAGKIRYIGVSNETAFGV
						MRYLHLADKHDLPRIVTIQNPY
				1		SLLNRSFEVGLAEVSQYEGVEL
						LAYSCLGFGTLTGKYLNGAKP
						AGARNTLFSRFTRYSGEQTQKA
						VAAYVDIARRHGLDPAQMALA
						FVRRQPFVASTL\LGATTMDQL
13729	44097	Α	13809	1	618	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	•	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13730	44098	A	13810	80	952	RPGIVALNAILPLSRTRQSKPTR LKKRAIANCAPTWKKRFVALV
	İ				ļ	YRTA*RFPSITLSVAVT*PASAK
		į				NKAIKMTQKIEQSQRQERVAA
						WNRRAECDLAAFQNSPKQTYQ AEKARDRKLCANLEEAIRRSGL
					İ	QDGMTVSFHHAFRGGDLTVN
						MVMDVIAKMGFKNLTLASSSL
					ŀ	SDCHAPLVEHIRQGVVTRIYTS
						GLRGPLAEEISRAYCADSHHGP
						FAMVNNMTDLTAQEPAWQTR
						DHLDDPVIGELRNRFGPDAFTV
						QATRTGVPVVWIKREQLLEVG
		İ				DFLKKLPKPYVMLFDLHGMDE
						RLRTHREGLPAADFSVFYHLISI
:						DRNRDIMLKVALAENDLHVPT
	}					FTKLFPNANWYERETWDLFGIT
	1.1000	<u> </u>	10011	00	706	FDGHPNLRWIP
13731	44099	A	13811	92	735	RLLRPLPQPRSSVLPGLTTFHAS
]			STSETRQSKPTRLKKRAIANCA
						PTWKKRFVALVYRTA*RFPSIT
		1				LSVAVT*PCVNIITIGCQVKTVT
						TIATAQIQRFTRFNHFSRFQYFR
						NSPKQTYQAEKARDRKLCANL
						EEAIRRSGLQDGMTVSFHHAFR
						GGDLTVNMVMDVIAKMGFKN
						LTLASSSLSDCHAPLVEHIRQG
						VVTRIYTSGLRGPLAEEISRGLL
		ł				AEPVQIHSHGGRVHLVQNGEL
						NIDVAFLGVPSCDEFGNANGYT GKACCGSLGYAIVDADNAKQV
						1
12722	44100	<u> </u>	12012	1	1113	VMLTEELLPYTDNPEA
13732	44100	A	13812 13813	5121	6228	LRRDSGRLRSCWHPRSPTSMGS
13733	44101	Α	13013	3121	0228	LGQREDLQDEDRNSGADPGQR
1						GRCSIHLEGPGQQPCLQRAVQG
						EGDPVLAKEEIQDQCHPHGQV
			j			QLLLVPAAEPVRAVPPRVQPVL
						PHHHHPAELSGIREE*GNQNDA
						EN*TISTTRTGSGLESSR*M/SILP
						LSRTRQSKPTRLKKRAIANCAP
						TWKKRFVALVYRTA*RFPSITL
ļ						I I
		Ī				SVAVT*PYWSGAPGGPCRRPYC HFRHRGARGSGAPGDGGEAPP
						AAAFLGGDVGFGL
12724	44100	1	12014	5.6	106	
13734	44102	A	13814	56	196	CRGDGGKLIIQCQRRMGI*PCG*
						LRQRKAELFQHLPLC*FWRPEI
10705	144100	P.	12017		2272	GTVCSGSIDNGSDRLRR
13735	44103	В	13815	64	2373	
13736	44104	A	13816	3	403	
13737	44105	В	13817	55	1040	<u></u>

SEQ ID	SEQ ID NO:		P .		l .	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13738	44106	A	13818	587	1533	VLLTAKFFPSMYSPIKTAGGPA
						ST*VL*MIISIDAEKAFDKIHQPF
						MLKALNKSENKIPRNPTYNGCE
						GPLQGELQTTAQGNKRGYKQT
						QQHDMHMSTKNQYHENGHTA
ļ.						QGMKGITQPSAFIPTAESNSFQP
						QVKTLPSPIDAKQQLQRKIQKK
						QQEQKLQSPLPGESAAKKSESA
						TSNGVTNLPNGNPSILSPQPIGIV
			į.			VAAVPSPIPVQRTRQLVTSPSP
	İ					MSSSDGKVLPLNVQSHQNCW
						WTRFNMSSLNVVKMTTISLTPS
						NSNTPLKHSASFISATGTTEESR
	1					SVPQIKNGSVVSLQSPGSRSSSA
						GGTSAVEVKVEPETSSDEHPVQ
,						CQENSDEAKAPQTPSALLGQKS
		1				NTDGALQKPSNEGVIEIKATKE
		1				RRSSPATEQSWMENVFEEQREE
						GFRRSNHSQIREDSQTKGKEEK
i						RLKELMELKTKVRELREECRSL
						RSRCDQLEERVSAMDDEMNEM
						KREGNFREKRIKRNEQSLQEIW
						DYVKRPNLRLIGVPESDGENGT
						KLENTLQDIIQENFPNLARQANI
13739	44107	1	13819	590	817	QIQEIQRTP
13740	44107	A	13820	107	410	KGRNIQLNGPSSRVSLWRSLYF
13/40	44100	108 A	13020	107	410	LNLNTGLPC*IGEVLLDNILQSV
						FQLGSILPITFRYTNQT*IWSFHI
						VPYFLEALLISFYSFFSKLPFSLH
						FIHFIFHR
13741	44109	В	13821	1	3735	
13742	44110	A	13822	922	1137	HFFLHFNFGESDNYVSWSFSSR
1.57.2		'	1.5022			GVSLWRSLYFLNLNVGLPC*IG
						EVLLDNILQSVFQLGSILHITFR
						YTNQT
13743	44111	В	13823	1	954	
13744	44112	Α	13824	3	156	
13745	44113	A	13825	683	972	SHRLLRLLHSSRSSPGLVSSSISS
						FKHFSVLVILFTLMIVSFAVQKL
						FSLIRSHLSILALVAIAFGVLVM
						KSLPTPMSSMVLSR/SFF*GFYG
						FRSYI
13746	44114	С	13826	1	895	
13747	44115	С	13827	372	1727	
13748	44116	Α	13828	705	1072	LALLVELIPLPLCNGLLCLF*SL
						LV*SLFYQRLGLQPLPFFVFHLL
						GRSSSILLF*AYV*LFWHDFAA
						AGTGCSFPCLALPSGALVGQA
						WW*QKSLSICLSVKDFISPSLM
		1	}			KLSLAGYEILG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	j			sequence		
13749	44117	A	13829	1	1136	
13750	44118	Α	13830	547	675	
13751	44119	В	13831	144	1877	
13752	44120	A	13832	1	441	
13753	44121	A	13833	27	344	
13754	44122	A	13834	287	600	GPGSLQSSPLLSPPWPRASSPLL
3,5,			1.505	207		LRPSVRGWHLGEP*NCFPSPLA
		1				AS/EQAPKRSS*PQCNHR*RETA
						ARGLAYPPDRPLPSMLSCESPE
						TPAKAGPGRERGCEKC
13755	44123	A	13835	2	226	II ARAGI GILIKOEEKE
13756	44124	A	13836	2	226	
13757	44125	В	13837	45	518	
L		В	13838	1	699	
13758	44126				362	
13759	44127	Α	13839	3		
13760	44128	A	13840	2	130	
13761	44129	A	13841	3	96	
13762	44130	A	13842	1	545	
13763	44131	A	13843	2	144	
13764	44132	Α	13844	2	91	
13765	44133	Α	13845	1	370	
13766	44134	С	13846	1	214	
13767	44135	A	13847	215	310	
13768	44136	A	13848	2	91	
13769	44137	A	13849	1	289	
13770	44138	Α	13850	1	997	
13771	44139	Α	13851	1	192	
13772	44140	Α	13852	2	267	
13773	44141	Α	13853	2	181 '	
13774	44142	A	13854	1	171	MESYAAIKNDEFMSFVGTWMK
						LETLILSKLSQGQK\TKHCMFSL
						TDVIARRRSWTSI
13775	44143	Α	13855	92	244	KRRRRRGQIRKMKEEEGNRRT
						WMKLETIILSKL*QRQKTKHRM
						FLLIGGN
13776	44144	Α	13856	146	382	
13777	44145	В	13857	124	1215	
13778	44146	В	13858	1	628	
13779	44147	Α	13859	377	484	
13780	44148	Ā	13860	2	220	WARTSCLKHQKQWQQKPKLA
				1		NGI*LN*RASAQQKKLPSEWTG
		İ				TWMKLETIILSKLLQGQKTKHR
						MFSLIGGN
13781	44149	A	13861	965	1150	GVDQRAVCSSRIW*SHLFLFHS
13781	'	 	1.5001			VVVHLPWAGTWMKLETIILSK
						LSQGQKTRHRMFSLIGGN
13782	44150	1	13862	254	571	YSYKLEIIIGKQPEIYRDPCYRP
13/82	44130	A	13002	234	J / 1	AV*SQTAVLAISEAPWA*DLPS
						7
				}		QVSDTISWWTVC*AHWKSAVL
						GWE*PGFPGTWMKLETVSLSK
L			<u> </u>	<u> </u>		LSQGQKTKHHMFSLIGGN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			İ	sequence		
13783	44151	A	13863	62	214	
13784	44152	Α	13864	274	435	
13785	44153	Α	13865	1	294	ASTAGVSYYVAQAGLKLLGLS
						LSKCRDYRCEPPCPE*MSLYKV
			1			MAMARKAMSL\YIYFFLDEFM
			1		}	YFAGTWMKLETIILSKLSQGQK
						TKHRMFSLVGGN
13786	44154	Α	13866	663	1266	PPLTCQCSHLCVTLVGH/WIKLE
						TIILSKLSQGQKTKHRVFSLIGG
	·		<u> </u>			AGALAKECMSLYPGCKITVFDI
						PEVVWTAKQHFSFQEEEQIDFQ
						EGDFFKDPLPEADLYILARVLH
	:				}	DWADGKCSHLLERIYHTCKPG
						GGILVIESLLDEDRRGPLLTQLY
				1		SLNMLVQTEGQERTPTHYHML
						LSSAGFRDFQFKKTGAIYDAIL
13787	44155	Α	13867	1	681	
13788	44156	Α	13868	1	247	
13789	44157	Α	13869	2	178	
13790	44158	В	13870	1	2042	
13791	44159	Α	13871	1	378	
13792	44160	Α	13872	1	683	
13793	44161	A	13873	1	972	
13794	44162	Α	13874	308	406	
13795	44163	С	13875	177	653	
13796	44164	Α	13876	1	465	
13797	44165	Α	13877	3	1175	FFSSQAAALLVVLGSSHTLQRA
						RKHSAPGLPDTCALLQPPAASA
						AAAPSMSGPDVETPSAIQICRIM
						RPDDANVAGNVHGGTILKMIE
						EAGAIISTRHCNSQNGERCVAA
						LARVERTDFLSPMCIGEVAHVS
						AEITYTSKHSVEVQVNVMSENI
						LTGAKKLTNKATLWYVPLSLK
			1			NVDKVLEVPPVVYSRQEQEEE
						GRKRYEAQKLERMETKWRNG
						DIVQPVLNPEPNTVSYSQSSLIH
]			LVGPSDCTLHGFVHGGVTMKL
			}			MDEVAGIVAARHCKTNIVTAS
						VDAINFHDKIRKGCVITISGRMT
						FT\SN\NSMRFKVLLDPAPVVDS
						FQKRYRGRQCLSST*VSLSQER
						QV/PLPVPQLVP\ETEDEKKRF\E
						EGKGRYLQMK\AKR\QGHAEPQ
13798	44166	Α	13878	121	361	SLFELENRSKFSVATAGLVPSR
						RLSINCIRRAFITSSNGFPLWFCL
						HQKGPVDCCVAVY*CGTEDGS
						WAPTHPGVGPSLP
13799	44167	Α	13879	1	462	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
E				sequence		
13800	44168	Α	13880	3	287	PCSLSSDCCASNQRDSVGIGPSE
						PRVGYNILVIRFLSPSEKRSPR\V
		1				GVTRFSSAWEPSQLAGILPISEP
						PSNRIFACWGKPAWTACCNSLR
						ARR
13801	44169	Α	13881	1	1332	
13802	44170	Α	13882	1	753	
13803	44171	Α	13883	1	516	· ····
13804	44172	Α	13884	71	277	IYFNTQESFCPPRPLK/LCDGRG
						CHKGL*NAFKAFSPFSWPLALG
						SFLLMQISAACLNSFPENGLFFS
						TT
13805	44173	В	13885	1	1095	
13806	44174	Α	13886	1	741	
13807	44175	Α	13887	264	2880	WLLQKHKLHVSR*LGL\PDTSW
						ILPSRPDAGGRFPIRTPPCVLVV
				ļ		HKPSRHCREWTANTVLPKGAL
			i			AKWLLEPKADLQGWHLPADG
						QYTKPSISYNIKLYINCYNRTVT
						TAAQHQQVQRKHGASIFLAEN
						VAVREKDKEERKVTRAPTEQA
						WTAGSWIETAYREIKGVGQES
						VCITVMHLEELVYELAWIIMSA
		1				VVEAFDPVAPSSKCRLNPDPDP
						YCCTPYTLHTSPPPPPSCCTPYT
						LHTGPPPQTPFCCTPYTLDTSPS
						PQTHSCCTPYTLHTVPTFHLPA
		1	}			AHPIHLHTIPPPQTQGQGFVSED
						EYLEISDIKRDQSGEYECSALND
						VAAPDVRKVKITVNYPPYISKA
						KNTGVSVGQKGILSCEASAVP
		ŀ				MAEFQWFKEETRLATGLDGMR
			•	}		IENKGRMSTLTFFNVSEKDYGN
			1			YTCVATNKLGNTNASITLYVD
						HIVGSKALLSKCKRTEIITNYLS
						DHSAIKLELRIKNLTQNHSTTW
						KLNNPLLNDYWVHNEMKAEIK
						MFFETNKNKDTTYQNLWDAFK
						AVCRGKFIALNAHKRKQERSKI
						DTLTSQLKELEKQEQTHSKANR
						RQEITKIRAELKEIETQKTLQKI
						NESRSWFFERINKIDRLLARLIK
						KKREKNQIDAIKNDKGDITTYP
						TEIQTTIREYYKHLYANKLENL
						EEMDKFLDTYTLPRLNQEEVES
						QNRPITGSEIVAIINSLPTKKSPG
L			<u> </u>	<u> </u>	l	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	I .	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13808	44176	A	13888	1	629	AMAGSPPPASLLPCI\LISDCCAS NQRDSVGVGPSEPGVGYSLVV RRFLSPSEKRSIRVGVTRFSRCR PSPLSLTQKGNSLTSWASQVLT PAPEKQPDNKLPLPRRGSRDPT EYFHGQEVRDKSDEHVDHVSQ IHSFRDEVKRLGAEDSSNAETP TELHEFNTGGVYRPPNSTSVIQP LDQGVIRTFKPCYTKYSLEKIVS AVEENPDRT
13809	44177	Α	13889	1	1452	
13810	44178	Α	13890	167	309	
13811	44179	A	13891	3	386	SVGALQKQQGWHQRRCISLHF PSFLCVFSLAKSRRADHNQDPR ASRMAFSLTSSPLARSAASAVA AFGCGTSGNGRFSSVP*AYRQV FQRIAIAVYGPLMRDR*AHPVK TPDAIDCRIRQPAIPRLQ
13812	44180	В	13892	121	298	4
13813	44181	Α	13893	473	1826	
13814	44182	Α	13894	1	270	
13815	44183	A	13895	3	3706	
13816	44184	Α	13896	1	1464	
13817	44185	В	13897	591	729	
13818	44186	Α	13898	1	1326	
13819	44187	Α	13899	1	1152	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13820	44188	Α	13900	1	2524	MREEFAHRWLARAGEKTPPDA
						LEKLTDPLMFAAAMVSAGKAD
						VCIAGNLSSTANVLRAGLRIIGL
						QPGCKTLSSIFLMLPQYSGPAL
						GFADCSVVPQPTAAQLADIALA
			,			SAETWRAITGEEPRVAMLSFSS
						NGSARHPCVANVQQATEIVRE
						RAPKLVVDGELQFDAAFVPEV
						AAQKAPASPLQGKANVMVFPS
						LEAGNIGYKIAQRLGGYRAVGP
						LIQGLAAPMHDLSPCIAATDAG
						SAAAQRIGDLVSVHVIPRPHVD
						LEQVFPIGLKAYTATCNHPVSL
						RFTIKSSSDRIKLSNQTLQSKNL
						FIMRFYCFSLCLSLSAFYHCRGV
						IVNIQAVFWLQKGTEAADILTT
						CAQLWIMMYALLSVFSLLDVIL
						NLAQKFPAASQLPLKGIFQGIKL
						IGAILVGILMISLLIGQSPAILISG
						LGAMAAVLMLVFKDPILGLVA
ļ						VRNWDNTITTIPTWSLVSDSFK
						NWSGMSASGGRRIKRSISIDVTS
						IRFLDEDEMQRLNKAHLLKPYL
		ŀ				TSRHQEINEWNRQQGSTESVLN
						LRRMTNIGTFRAYLNEYLRNHP
						RIRKDMTLMVRQLAPGDNGLP
						LEIYAFTNTVVWLEYESIQADIF
						DHIFAIVEEFGLRLHQSPTGNDI
						RSLAGVTINKAPGDDEILKNTV
						DFVSFSYYASRCASAEMNANN
						SSAANVVKSLRNPYLQVSDWG
						WGIDPLGLRITMNMMYDRYQK
13821	44189	A	13901	1	2370	
13822	44190	A	13902	1	282	
13823	44191	Α	13903	1	714	
13824	44192	A	13904	3	506	
13825	44193	A	13905	459	551	
13826	44194	A	13906	401	492	
13827	44195	В	13907	479	702	
13828	44196	A	13908	120	256	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13829	44197	A	13909	16	1249	SKSGAIEIARLAGRVHGVVVQI
						TSEIGPLTLLTANFAFSAS*STA
						WKATSMDGEVLS*YSTSASAS
						AERQSTHQ*TGFAPLCR*PLPMI
						/VAQRADDVGFSFEVHGQVRV
						RPVAQHAQTDKVFALTVNLGR
						RVFAALGAELGGARSAVRPVP
						ECYAVRETLALLQKIFPIRQCEN
						SVYRNRSRPCLQYQIGRCLGPC
						VEGLVSEEEYAQQVEYVRLFLS
						GKDDQVLTQLISRMETASQNLE
						FEEAARIRDQIQAVRRVTEKQF
						VSNTGDDLDVIGVAFDAGMAC
						VHVLFIRQGKVLGSRSYFPKVP
						GGTELSEVVETFVGQFYLQGSQ
						MRTLPGEILLDFNLSDKTLLAD
						SLSELAGRQINVQTKPRGDRAR
						YLKLARTYAATALTSNFRSIYR
						SPATDRACQRVETAGNER
13830	44198	A	13910	1	438	
13831	44199	A	13911	1	1200	
13832	44200	A	13912	1	300	
13833	44201	A	13913	1	431	
13834	44202	A	13914	2	1465	
13835	44203	A	13915	1	513	
13836	44204	Α	13916	3	535	RDKILVFKDENFWMIRGYAVLP
						DYPKSIHTLGFPGRVKKIDAAV
						CDKTTRKTYFFVGIWCWRFDE
				Ì	ļ	MTQTM\EKGFPQRVVKHFPGISI
						RVDAAFQYKGFFFFSRGSKQFE
						YDIKTKNITRIMRTNTWFQCKE
						PKNSSFGFDINKEKAHSGGIKIL
	11 (FFI)					YHKSLSLFIFGIVHLLKNTSIYQ
13837	44205	Α	13917	1	876	
13838	44206	Α	13918	23	414	
13839	44207	Α	13919	2	425	QERGPHQVIQRKECLREYSPVV
						S*AKGTGVAVGQKGTLQCEAS
						AVSSAEFHWYKDDKRLIEGNK
						GVKVENRPFLSKLIFFNVSEHD
						YGNYSCVASNKLGHTNASIML
						FGPGAVSEVSNGTSRRAGCVW
						LLPLLVLRLLLKF
13840	44208	A	13920	1025	1270	
13841	44209	A	13921	370	701	
13842	44210	В	13922	1	660	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13843	44211	A	13923	<u> </u>	432	RFSRVDDFVLTMDDGPPPRRAR
130.5		`	1.0720			GDLTHSGLWRVCCIEGIYKGHC
						FRINHFPEDNDYDHDSSEYLLRI
						VRASSVFPILSTILLLLGGLCIGA
						GRIYSRKNNIVLSAGILFVAAGL
						SNIVGIIVYISSNTGDPSDKRDE
						DK*NHYN
13844	44212	A	13924	1	1093	
13845	44213	Α	13925	1	1203	
13846	44214	A	13926	1	1353	
13847	44215	A	13927	2	445	
13848	44216	Α	13928	1	227	
13849	44217	Α	13929	1	840	
13850	44218	С	13930	120	356	
13851	44219	Α	13931	40	342	
13852	44220	В	13932	403	473	
13853	44221	A	13933	70	2169	DOA A TOOMOGED VODEN/DELITIV
13854	44222	Α	13934	1428	1905	RCAATSGYGERVQRTV/PTHTV
						W*ASLSQSSAHRVVAINSIANK
1	:					RSSMPPKSWATVALTSTKLKCR
						FMAKVRNKSFSVLHNV*MTISI/
		l				SPWVILSSGVDEKLFPRAVRVA
		ŧ				MTAGASGFLAGRAVWASVVG
						LPDNELMLRDVCAPKLQQLGDI VDEMMAKRR
13855	44223	A	13935	227	342	VHSWQRYSGR*RGY*PQNQSA
13833	44223	A	13933	221	342	TDQTRRWQSLKTAGAVA
13856	44224	A	13936	480	928	FFADLQQPVCADSYPLLVQMK
13630	44224	^	13930	1400	720	LAPHKS*NAVLWQRSATKSFSV
						L/AQGLNDHINMPWVILSSGVD
						EKLFPRAVRVAMTAGASGFLA
						GRAVWASVVGLPDNELMLRD
						VCAPKLQQLGDIVDEMMAKRR
]			FIPLLRRWVDLALTRWLITVSK
13857	44225	A	13937	564	1185	WLHASAALLRW*YQLPGQPHH
,		1				RRCRSADGSHVPSLCAAVPDQR
						\PLITVRRIGMQAMTTSTTGDRQ
						RSWQIAMDGSQKLPQRMLDSV
						RWHLAHDSKFDLLALGVAGW
						MRYVGGVDEQGNPIEISDPLLN
						RMAANRFVHITOLPVIYRAFRN
						SFTQVARLQLCHKRFQQRHTSV
						REQIVTVLSGIRDNHVHFVQTL
						TGDGVGNQRQLVQ
13858	44226	A	13938	228	424	HRFCPAARH/RSWSGLREMAVS
						ISMPRRIINSGRYCAICWYKRRS
						RCARNMLIFPVARSTPIPGPS
13859	44227	В	13939	128	1670	
13860	44228	Α	13940	2	2367	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13861	44229	I A	13941	1	921	ELPASPAPCARTPQPLGGRWEA
13001	1722	` `	135,11	Ĭ,	1	ARRKERVLCLFDVDGTLTPAR/
						QGRTLKIGSSHRECHADTPEAE
			ŀ			CGVCSSVTEVRAKWRSQTC*KP
						ECYASPLPP*KIDPEVAAFLQKL
						RSSYLPVGSQTWSHTLSSSCPLC
						DPEVIGRDTVPVSGSGSPGHAQ
						SPVFLGQTIQNHLGPGGARFEF
						RA/CSGSLWLPALPHH\RGTFIEF
						RNGNCVPSQGLAHSRSSELPSA
		•		-		PSPLQKEQAALLLALSAPPSRE*
]		QSPPLPTGGMISFDVFPEGWDK
						QVRSSQCGEGLHLNIPPLPILCV
			1]		QGGNDFEIFADPRTVGHS
13862	44230	A	13942	105	416	HSVLEVGPWWEVFGSWGQILH
						SLVSSHHVRHWLLPSPSAMTES
						SLTFQQKPSRCQHHASCKAYHS
						HQLLLVPTAGRLEDRSHH*IPC
						KHSPAPAWSLAAPVGN
13863	44231	Α	13943	191	573	SSSGGYQQLMRVIGTRDTAV
						NKASNVPSFGA*I/PSGRGQQAK
						LIALTRALTLAKELHVNIYADC
						KYAFHILHQHDVIWAERGFLTV
			ļ			QGSSSIINASLIKTLLKATLLPKE
						AGVIHYKGHQKASAPIA
13864	44232	Α	13944	1	1065	
13865	44233	A	13945	321	537	
13866	44234	A	13946	346	1260	GGPARAEFSRHLGCKGHHPLPK
						DVASEVPAGSSTSVSSLIRTNGC
						PGLGSWEPP*FFSENKRLSTLMS
						FTLRDSKMFSPQSGLFLGCSAG
					•	EHSALRDPTASGLPGTASGVRP
						GR/RQRVGSDIEASTQGESKPCI
						LSI*TLPNA*LKTKTLTPLEIAAC
						TNPHSFSITARIGPAHLAAPRPV
ł						TPAGRGGARAPGR*CAAARPPP
						ASQ*VSGSPAPGGGPSYAPAAR
						TS*PSSLAPGAKRLRFSRVATFC
						CSFLAWSSSSNKAFCT*EHMSL
				•		LSSWISQSLEKVASCCPCEQFDS
12967	14225	 	12047	1	459	WKRLENILLFS
13867	44235	В	13947] 1	433	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	, , ,	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13868	44236	Α	13948	63	1263	VEAAKKAHHAACKEEKLAISR
						EANSKADPSLRPQAHTYALSPL
						FLTAMPSV*QGEHVIYTRVRHS
						DQVSESSGKPLSLPLLEPGIWAF
						EAAQVPPRWDQVRSLEPTTEW
						EAPGLKKRRVSQGLNGNEPLSC
į						AWIFL*TKEKYEKSLKELDQGT
						PQYMENMEQVFEQCQQFEEKR
						LGWDKG*LAPAF*KSHVSL*FS
ŀ						YKAIYHDLEQSIRAADAVE/DR
						RGF\RK**TFGFSPSAPVA*EWS
						ADLNRTLSRREKKKATDCQDP
						HGPLQSWDPLKALSLHSTLNVP
						SNPAQSAQSQSSYNPRGPHTLC
						RPVLRALWPPLSCSVSSYEKTQ
	<u> </u>					SYPTDWSDDESNNPFSSTDANG
i						DSNPFDDDATSGTEVLCSPPSV
						SYLSWRRLPLCLAGDELTKME
						DEDEQGWCK
13869	44237	Α	13949	2	485	AAAFGTRLSRRGRGCGVTGSG
ļ						GGRRRVPAPRRPRSPLQPPHQ/P
						PRRPSRRTPGCGSRGA\PSRGRR
						LVPPPPEPPRAAQQRARH\HRTS
						\RPAARQPPRRPPAQPRSCPRRV
						RPTPAWSPPGRRGPRRSRRRRR
						RGGRARARRPRRRGQVPRPQS
						ATGAGHAR
13870	44238	A	13950	1	207	
13871	44239	В	13951	1	5297	
13872	44240	В	13952	1206	1360	
13873	44241	Α	13953	46	369	NEGKNAKGSQRERSGYPQREA
						HQTNSGSLGRNPTSQKRVGANI
			:			QHS*RKEFSTQHFISSQTKLHK*
						RRNKILYRQANAERFCHHQEGS
100=:	11010		10051	17.6	260	TKHGKEQPVPAAAKSCQNVK
13874	44242	Α	13954	476	868	KYDCCISSTFCRALVHFSRTTFQ
]						VNSPLSPNISAFGYMPLKYSIYF
						LHHLRPNIL*AASYFLKYKGTL
1			·			KLPALRGCLS*STRCWCKIHICI
						*ND/ECLQFLGPWFENEFCFLPS
L			<u> </u>			TSQFVEFVLDILVLQNEP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13875	44243	A	13955	11770	3003	PANIIMTGS\NSHITILTLNVNG\
100.0			10200			LNAPI*RHRPAIA/S/WLKSQDPS
						VV\CIQETHLTCRDTHRLEIKG
						WRNIYQANGKQKKAGVAILVS
						DKTDFKPTKIKRDKEGHYIMAK
						GSIQEEELTILNIYAPNTGAPRFI
						KQVLSDLQRDLDAHTIIMGDFN
						TPLSTLDRSTRQKVNKDIQELD
						SALHQADLIDIYRTLHPKSTEYT
						FFSAPHHTYSKTDHIVGSKALL
:						SKCKRTETITNCLSDHSAIKLEL
						RIKKLTQNHSATWKLNSLLLND
		ļ				YWVHNKMKAEIKMFFETNENK
						DTAYQNLWDTFKAVCRGKFIA
						LNAHKRKQERSKIDILTSQLKE
						LKKQEQTNSKATRRQEITKIRA
				,		ELKEIETQKTLQKKQPNKDPAM
						EKNNKTDRVLARLIKKKREKN
Ė						EFSAHILYKFSKQNGS
13876	44244	Α	13956	105	392	RSRKPRGNPKSHPEDTDPTKAP
						RKRQISSRGHRSHR*EPGPQSAA
		İ				PGFREQRRRGQMG*IWYLTPTP
						PGGQRPSSAASPRGFPPTNNSSR
ļ						LPAEPE
13877	44245	Α	13957	55	403	SEGMSGKASEKQEAFACPRHLT
						RAVDGPS*SGLLGSRLPPSCSPG
			<u>.</u>			KEAPGGSELYLCLMRGLQTL*A
}						RSGTGLR/PNPAEIR*EPGPQSA
•						APGFREQRRRGQMG*IWYLTPT
						PPGGQ
13878	44246	Α	13958	217	444	RSRKPRGNPKSHPEDTDPTKAP
		İ				RKRQISSRGHQSHR*EPGPQSA
						APGFREQRRRGQMG*I*TPCRP
						ALPKSKPRPL
13879	44247	A	13959	309	670	PCPRASSQGLKATH\GQR*SSEP
]				PGASFPGLQDGGRREPRRPD*D
		1				GPSTGLVRCLGHANVGDPCLK
						VFVFFKNLSPTYLRASAEQTLP
						LLLPHLHGLCLHQPLHLGFTAA
						WAQLTFWEASQ
13880	44248	C	13960	392	760	

SEQ ID		Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
•				sequence		
13881	44249	Α	13961	24	915	HAGRHARGSLFPGARPSETRAH
1	-					CVRTARPNCKEEDTPMVRGKN
						SGGEHCTWGPSLAAGG*GIGDP
		l				GGLPGT\PHPGLQLLLHGFAGP
		l				HMASLCCGVSSPAGVRQWPA*
						GSGNHQVKSWLGLPKNVS*AP
						SML*NPKMQRLMKAQPHGRCE
						VRRGRVCSALGP/IGMSGKASE
						KQEA*DEGHPHLHAPGT*QDQ
	-					VDGPS*SGLLGSRLPPSCSPGKE
						APGGSELYLCLMRGLQTL*ARS
						GTGLRANPAEIR*EPGPQSAAP
1						GFREQRRRGQMG*IWYLTPTPP
						GGQRPSSAASPRGFPLV
13882	44250	A	13962	397	612	RPQHHTFCCDGRVMMARQKGI
						FYLTLFLILGTCTLFFAFE/CSFG
						*VELPDYGYFGSRGRTCFLGSL
						GLVLG
13883	44251	Α	13963	125	1298	FQSWAQPLFLLSCNRKTHFGTR
						IPI\MSVMVVRKKVTRKWEKLP
		1	İ			GRNTFCCDGRGMMARQKGIF\
						YLTLFLILGTCTLFFAFE/CGRYL
						AVQLSPAIPVFAAMLFLFSM\AT
		İ				L\LRTSFSDPGVIPRALPDEAAFI
		1				EMEIEATNGAVPQGQRPPPRIK
						NFQINNQIVKLKYWYTCKIFRA
						SRASHCSICDNCVERFDHHCPW
						VGNCVGKRNYRYFYLFILSLSL
						LTIYVFAFNIVYVALKSLKIGFL
						ETLKETPGTVLEVLICFFTLWSV
						VGLTGFHTFLVALNQTTNEDIK GSWTGKNRVQNPYSHGNIVKN
					•	CCEVLCGPLPPSVLDRRGILPLE
			!			ESGSRPPSTQETSSSLLPQSPAPT
						EHLNSNEMPEDSSTPEEMPPPEP
						PEPPQEAAEAEK
13884	44252	Α	13964	44	647	DYSAQHGKASQKHVARLISSGT
13004	744232	^	13304	 	047	FQGATVSRPFQLSTSSSSFCSSSS
						SLSSSSSLSSSLLLSCSSESVFLS
		-				LSFFFCSEASFLPFCSRLYASRA
						SFNGVTNRWNSISSMAESTSLA
						FSVFRFPFIAKLLAQDVAYSTKT
						LAAREIGAPGLLGDVDTVRELR
}						GDRG*AKLPFQHLLPSCPFVRPP
						RAAHRLLDDPGDHGIGQV
13885	44253	В	13965	158	1075	id i i i i i i i i i i i i i i i i i i
12002	77433	٦_	13703	130	1.073	L

			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13886	44254	A	13966	125	1297	FQSWAQPLFLLSCNRKTHFGTR
		1				IPI\MSVMVVRKKVTRKWEKLP
						GRNTFCCDGRGMMARQKGIF\
						YLTLFLILGTCTLFFAFE/CGRYL
		ı				AVQLSPAIPVFAAMLFLFSM\AT
		1				L\LRTSFSDPGVIPRALPDEAAFI
						EMEIEATNGAVPQGQRPPPRIK
						NFQINNQIVKLKYWYTCKIFRA
						SRASHCSICDNCVERFDHHCPW
						VGNCVGKRNYRYFYLFILSLSL
						LTIYVFAFNIVYVALKSLKIGFL
						ETLKETPGTVLEVLICFFTLWSV
			•			VGLTGFHTFLVALNQTTNEDIK
						GSWTGKNRVQNPYSHGNIVKN
						CCEVLCGPLPPSVLDRRGILPLE
]		ESGSRPPSTQETSSSLLPQSPAPT
]		EHLNSNEMPEDSSTPEEMPPPEP
						PEPPQEAAEAEK
13887	44255	Α	13967	247	850	DYSAQHGKASQKHVARLISSGT
		1				FQGATVSRPFQLSTSSSSFCSSSS
						SLSSSSSLSSSLLLSCSSESVFLS
						LSFFFCSEASFLPFCSRLYASRA
						SFNGVTNRWNSISSMAESTSLA
						FSVFRFPFIAKLLAQDVAYSTKT
						LAAREIGAPGLLGDVDTVRELR
						GDRG*AKLPFQHLLPSCPFVRPP
		ļ				RAAHRLLDDPGDHGIGQV
13888	44256	В	13968	298	380	OR A CONVICENCY OF BUY BUY BUY
13889	44257	A	13969	48	252	QRAGSPHSPRSLSAPPLPELPLW
						RHLRSPSAHRCTVGAPFWDGQ
		ļ				G/LEPAPSAALAAFPRSRARDLQ
12900	44260	l _D	13970	1	447	LA
13890 13891	44258 44259	B A	13970	236	559	MWLEPMQMGFLHMMEKMAA
13691	44239	A	139/1	230	339	RTSAILD*GTLK*FHFTLTTSLK
						ALSSHTPIFPGTGELQLPVSPSV
	1					CLDQGMQLKPSTSSHLLKTVKP
						RMKRQSLLHMKQSFEPKIYL
13892	44260	A	13972	1	645	MKPQTLTVSVTAFKVAHLEFVP
13072	44200	^	13972	1	043	SDVQMCSEFLPSGGFVVSLASG
						VKLQTFAVSFTAHKSSVDPKNS
						EAQLASPSGFCTRAADGAACQS
						\LPCTCTPQPLGGRWDWALWS
						RGW/LLVEEAWAAQEPTEELRE
						RALALASPERGSHSAAVGQRAP
						QVPPKWEPRQRRRRERARAVR
						TASTLSPLNPPSKQDTLTAVGN
						LADDHSSYFLLDRGEEGALQL
13893	44261	HA-	13973	28	322	Z. I.D.D. I. D.D. I. G. D. C.
			<u>. </u>			
		1	13975	942	3414	<u> </u>
13893 13894	44261 44262	A	13973 13974	28 1760	322 4426	\LPCTCTPQPLGGRW RGW/LLVEEAWAAQ RALALASPERGSHSA QVPPKWEPRQRRRR TASTLSPLNPPSKQD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	· · · ·
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13896	44264	Α	13976	1	274	
13897	44265	Α	13977	545	1150	RISGPGAVLLPVMPALWRLRHR
						MFSLTKK*HLPAHRSLAEGYLL
						YQPQLDTLKARLTQKYQELQV
						LFEAYQIKKTKLDRQSSSASLET
						LLALLQAEGAKIEEDTENMAEK
						FLDGELPLDSFIDVYQSKRKLA
						HMRRVKIEKLQEMVLKGQRL/S
						TRPWPRCPPGCPNWHLPPPFPT
						LPQSQWASCRCTSAHPPPTTPG
						AAGRL
13898	44266	В	13978	213	611	
13899	44267	Α	13979	1	639	MEAVTVVPVGTEEGMGEEEQD
						SGTTTGCGLPSVEKMLATNPGK
						TPISLLQKYGTRIGK\TPVYYLL
						KAEG\HQPNFTFWV\VGNTSCT
						AQGPRKKVVKHKAAEVALRHL
l						KVESMLEPAPEDSSSFSPLDSSL
						PEDFPVFTAAAAATPVSSVFLT
						RSTPMEKQPPLLP\QQSECIPVG
						ALQKLVVQKGWWLPEHTVT*E
	•					SRPDQHEELT*RVERFT
13900	44268	Α	13980	361	894	AFFPMSTAGVCGNVEGEPETP
						WSLPLSQVPFSPNA/CPQGTLGR
ĺ						S*AGVAGAPGPGVPGRSPAA\GI
						PEPAARPGRPALSRLPGLGGAG
						ARSGAA*G*GSRPGG/PLPAADP
						VQSWGP*EAPSHP*PGT\SADSP
•						PSGVSGATGDCLWGSGRAELF
				<u> </u>		GFATTSSAAPFPGLPTFFFIPVSP
13901	44269	В	13981	1	3842	
13902	44270	Α	13982	3	407	DAWAAARPGRSCALPPPGA/PE
-						EPGHVPGAA*G*GSRPGGRFQL
						PDPVQSWGP*EAPSHP*PRNFPQ
						TRRHLGVKQHGCSQPSPHGSRA
						SQAHPGISGLF/GPHHPGPPLFIH
]						GPRCCCAQATASNHLFACGTSS
		<u> </u>				DP
13903	44271	C	13983	127	297	
13904	44272	A	13984	3	1920	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	1	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13905	44273	A	13985	3	2271	GQGRRTAGGGGGGGGAGVGAG
,						AGAMEAERGPERRPAERSSPGO
						TPEEGAQALAEFAALHVWSGE
				ļ		GIGVGSPAHTPADGNSGPQVFD
						AGEVFGIMQVEEVEEEEDEAAR
						EVRKQQPNPLAPPTSLCPMPTR
						QPHIVACARSIFLIDHAWTCRV
						EHARQQLQQVPGLLHRMANL
						MGIEFHGELPIWVPEGSSGGW
						MGEREVPYVLCLQTAEEKMPV
						WYIMDEFGSRIQHADVPSFATA
		i				\PLEEGPQPSVPSPPLMGTH/DL
		ļ				GTEEVTRDFAYGETDPLIRKCM
Ì		1				LPLSSSHRGMETPCVHTSRASL
}		ł				PLQACELPSWSLLLINEPPRPAF
]				PSLP*PHLYFRCVSF*AIPGLVLE
						SRLDTWEPGSPSSGSPHRVYTD
						VQQVASSLTHPALPVRASRHCR
						PPLELAHLPPPCRKLSQERPGVL
						LNQFPCENLLTVKDCLASIARR
						AGGPEGPPWWVLGWHHLGTQ
						APGAPQTALLLPRGEDNHWICK
						PWNLGLCVLGSPRGLCPGHQR
						ACTCVLQVVSKYIESPVLFLRE
						DVGKVKFDIRYIVGGTSLAGTR
						PLKNPYVSAHLSSRAFAQGRRG
						RGGPTWSGWLLARDPARV*DL
						LSSLVHSVERCPMQWALSEWQ
						LHHLSTWSSGLCGLAG**VFQM
		1				HRTPTPP**QAACSRPSGGLYR
	1	l				A*WALCLLPGAADPLLRGCSEQ
						AS*LRLPGPQAEIFRAFTELFQV
13906	44274	Α	13986	1	600	MEEQRVQFTLWLHFSPHGICQF
						ESGRQEAKKLKGSGQPLKTGEI
]						KTGIQKYRREALLQHYLGPASL
						AGRSKALTGTAERSCSSGSDSSI
1						FTLCVNYNDSRAIMKTLAAIGT
						GFDCASKAEIQLVQILGMPPER
						FMYANSCKQMSQIKYTANNGV
						QMMIFGNEVKLMSVARTHSKQ
						P/CHISVKSGAMLKASRLLLE*A
	1				20.40	KEL
13907	44275	В	13987	1	2040	LARL EMONEONOUR COOCTOOL
13908	44276	Α	13988	205	438	LYFLETGVFCVQVPAQQC*CSV
						PGSCRRPLEEAGWRGLGLSGRL
						AGTGLPSGSRGNRTERGERGKL
	<u> </u>	<u> </u>	<u></u>		<u> </u>	VRGRRGVTRWLS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13909	44277	Α	13989	284	632	KTGVFCVPGPRPTVLM*CSFVG
						GKKNQYCLTGDAGCLQEKHSL
						GPAGV/HLEEAGWRGLGLSGRL
						AGTGLPSGSRGNRTERGERGKL
						VRGRRGVTRWLSHPQVQRQAD
						AGTCSPFEP
13910	44278	Α	13990	594	1003	VRKGHHAVRGLCGRAQFRCIS
İ						GDPGKRPLG/LPGGNAPKSGSL
		l				ARRPQPVRIPGSRPQ/RGSVSHL
1						GPAGVH*R/RAGWRGLGLSGRL
						AGTGLPSGSRGNRTERGERGKL
						VRGRRGVTRWLSHPQVQRQAD
						AGTCSPFEP
13911	44279	Α	13991	1	1500	VRVAHIPIAAAEIAAIPDRGGHL
						SVGKRKLTFWFFYGVLPFRDV
						AIEFSPEEWECLDSAQQHLHRD
						AMLENYGNLVSLAAGDQSLPG
						DAAASAVCRQLSPSACGAILGL
		l				PPVNGVAPVRLRGEIVTYTWSQ
		l				LTGLALALSKDGADGLRRLTRP
						TRRAPPALLLAPQTAAERWRG
		i				KCDPRHGGGCALALQGWTLYP
						RESPSRESKELGGLWSFRADLD
		ŀ				SRRQGFEEQWYPRPLRELRGRG
						GRRPEARRWEGGSPGPCRLLPL
						SAPELAPYRKLKVNGLRGAGL
						GDGETGGTGEGRGGRRRKRCC
						RTLSRDEGFNDICQDWWLRQF
		l				VGWVLYEQEVTLPEQWTQHLR
						TRVVLRIASAHSYATVVSAARS
						RQGRGIGGLIYTLGMPFFLSGW
						QVPEPHAAGYVRECLIAGPLLM
						P/TTCCPLLISPVSAVGEWGRHA
						RA*GADSSSLVQNTDFDFFNYA
						GLQR\SVLLYTTLTTYIDDITITT
		<u> </u>				GVEHDSAEIPIFPEYLRVNTGN
13912	44280	Α	13992	2	307	
13913	44281	Α	13993	220	427	MSTLPFPSLQMFRRPPYGSPSTC
[RSQDNRSYSRVTEELQERRRW
						NSGDAN*DQSPEARRQRPRHIE
	<u> </u>					EAA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1 2 2	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13914	44282	Α	13994	1	2832	MGKYYTQGDKVLMPLAIQVH
13711	1.1202	-				HAVCDGFHVGRMLNELQQYC
						DEWQGGAYPKGYFVQNTDFDF
						FNYAGLQRSVLLYTTPTTYIDDI
						TITTGVEHDSAAAPMVDSLIAR
						VGVMARGNAITLPVCGRDVKF
						TLEVLRGDSVEKTSRVWSGNE
						RDQELLTEDALDDLIPSFLLTGQ
	Ì					QTPAFGRRVSGVIEIADGSRRR
				1		KAAALTESDYRVLVGELDDEQ
						MAALSRLGNDYRPTSAYERGQ
						RYASRLQNEFAGNISALADA
13915	44283	Α	13995	1	2934	MITFDFMSHIQVTLMQEVSFHG
13913	14203		13773	[*	2754	LGQLRPCGFAGYSPPPGCFHRL
į						ASSVCSFSRCTVQAAVDLPFWG
						LEGTSPLLAASLGSAPVGTLCG
		1				GSDPTFPSWTALADVLHEEGHT
l						PASNFCLRNLAGEASQSWRKE
			:			GLGTPRPTASSTEELQTLNEDSR
						LMTSPHNLEETRDCDTELPKKS
						LGWKEIRVYGQNKTLGFWLQA
1	1			1		QLLICDGGNLRMFRLLPYGSPS
						TCRSQDNRGYGGVTEELMERK
		l				RWNSTDGFHIQGVLE
13916	44284	Α	13996	68	425	SHILPGAPGAPAWWTRWPSTLP
13710	144204	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	13770		1723	EPFPRGRGSPAGTSPISRPGLVQ
	1					SS*ASRGSDSRLPV/GPASCQAS
ŀ						GPGPDSRRPPPCTPA\GPHHGSL
						PSAGRVGASAAAAGPPSPAVPL
		ŀ				PPAERPAP
13917	44285	A	13997	324	723	TAFSLNFQGGVEGEVWAGTRA
13917	44203	A	13991	324	/23	ARGACAGERGLGGSRTRSGW/P
						GTAGPGQAGHGTCSLPCPSLPP
						AMGSCVSEPPRQAPPPAPWRLV
						PSTAQGLRSVGTQCGTGGSSAC
						GPC\GDPGF*THQSAPCQNGPIS
						SL SL
12018	14296	<u> </u>	14000	39	383	OL.
13918	44286	Α	J14000	الادا	دودا	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13919	44287	A	14001	181	1446	EDLLQEAQALAEVTAQGKGLG QVRSLDLSLEACAGSPPLQVLD LQDREPRSLTSLPSSAANRGKL EKAQSNPCGHRRHQAAGLKLP LFSAQAPQGQPTPAPRFRAATR KPGTAQSHAPLWSPAMSFFPEL YFNVDNGYLEGLVRGLK/ALGE LSQADYLNLVQCETLEDLKLHL QSTDYGNFLANEASPLTVSV\ID DRLKE\KMV\VEFRPHEGTMPY EPTSPSFP*TSITLQFT*FDNVI/L WLIHRARLHQRSIAELVPQVAH PLG\SFEQMEAVNIAQTPAELY NAILVDTPLAAFFQDCISEQDLD EMNIEIIRNTLYKAYLESFYKFC TLLGGTTADAMCPI\LEFEA\NR RAFINT\INFFG\TKTVPKKDRCP AFFPH/CGAGFFPEGPAAMGFG LNNLLTGAKKRGPYNFPEVQL AFLRGDG
13920	44288	A	14002	1	1182	
13921	44289	A	14003	26	342	ARFLAGP*LPSRRTGLRDLQPA MPEPP\PTPWAPVRPEPPRRAPP PAPRRPVPSTGKEQTTQGLRSA GAGHRDWQAAPPAAPVRDPLG EASWAPESGGDVESLYV
13922	44290	Α	14004	66	463	
13923	44291	Α	14005	3	1446	
13924	44292	A	14006	221	852	NSFLMVGFFPFFSL/CAAFDTKT GLRVAVKKLSRPFLRPDGYYTE FNIYLLFNFYILTSSAV*VVLSH SVVIECLLSIRHCSRYFRKSSED KNFLRGSRLLCNFSLISRYLVTH LMGADLNNIV*RYR*YK**FFK MNSPFLPLPVGGELIMHHKVDP VCFVFFSLPVSKS*QFVCSSSAL *ILDFGLARHTDDEMTAKGQPE CFVRWACAS
13925	44293	A	14007	55	784	RHIQDPASQRLTWNKSPKSVLV IKKMRDASLLQPFKELCTHLME ARGAGGGHDARSLTTAPVSRE PPCRRVGSAGGMSRLALCFQEN MIVYVEKKVLEDPAIASDESFG AVKKKFCTFREDYDDISNQIDFI ICLGGDGTLLYASSLF/QGNAA VVLRSRLKVRVVKELRGKKTA VHNGLGEKGSQAAGLDMDVG KQAMQYQVLNEVVIDRGPSSY LSNVDVYLDGHLITTVQGDGVI VSTPTG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
ļ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	ı			sequence		
12026	14004	<u> </u>	114000	1125	1502	DECRISORED DE ANELTEMENTES
13926	44294	A	14008	155	1592	PESRHQCFSDRSANFLTMEMEQ
						EKMTMNKELSPDAAAYCCSAC
						H\GDETWSYNHPIRGRAKSRSL
1						SASPALGSTKEIRRTRSLHGPCP
						VTTFGPKACVLQNPQT\IMHIQD
						PASQRLHVEQRPPRTVLVIKKM
						RDASLLQPFKELCTHLMEENMI
						VYVEKKVLEDPAIASDE\SFGA
						VKKKFCTFREDYDDISNQIDFII
						CLGVDGTLLYASSLFPGQPSPP
						VMAFHLGSLGFL\TPISFENFQS
						QVTQVIEGNASCCSSGSRLKGQ
	1					GW*RSLRGKKTA\VHNGLGEK
						G\SQACRPGTMDVGKA/QAMQ
						VPRS*NEV\VIDRGPSSYLSNVD
						VYLDGHLITTVQGDGKARSTV
						LGP*ASLGRE*RLRLSLSGVIVS
		1				TPTGSTAYAAAAGASMIHPNVP
						AIMITPISRSVLPTAGGPHLSLL
						DPQIMLSPEARNAGGGPFWKP
		1				WLRLLLSRPSISITTSCYPLPSIC
						VRDPVSDWFESLAQCLHWN
13927	44295	A	14009	327	466	
13928	44296	A	14010	2	196	
13929	44297	Α	14011	923	1776	WRKCGCETAGSASATRAAATE
						KMEAPRNKMCRDQENSAWRN
						AVPGP*E*RQPEEFPRTTSILAEP
						TVLSRRKCRPSEKRGSATEKFG
						ATSAVTENPPLGAAGERANKTL
						GAATVTPTLGPRPRMEKGGTR
						CPSTL*PSVQERQPESPSGNHPE
						TQKIEFAGRRVTFQHRWATTLR
						TRISGTGASPRDRSAFFGFPWC
						G/AAQPSDEESTPATQEEAQAQ
						AGVAAAASSEEPGHCAPRPPHA
						PRSALFEDWSREPERSDDGDLS
						ECRRPATEPWRPPGSAWRLPCA
						A
13930	44298	A	14012	568	1223	SARGLLPGSGPLSHAGPQLQVE
						WFGCLPPPDSTLELVVCGTVAI
		1]			SRVRMSAVCVLRTWS\RNAG\Q
						LI\CFTVFQPCSNVHVL/KGPNY
		1			1	VCFFG\YPSF\KYSHP\HNFLKTI
		1		į		NAVCGQ\LVQFRFPDTEEGIRK
						VTV/KCYVKEGDTVSQFDNIGK
				-	1	VQSDKASVT\IASPYDGVI/RKL*
						HSLDDIAYVGKPLVNIETEALK
		1				GTVNLFYQIDYCPPFVIGSQLK
L		İ	l	<u> </u>	1	OT ANTEL LATIN LCLLL AIGSAFY

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13931	44299	Α	14013	5	296	LFSRVPSKGPPRYHHTSSRG*TR AD*LISS\GNS*AHCIPSGLQGM LNLLPTPRAPVRPEPPRRAPPPA PRRPVPFDHPRACGTGRQLHLQ PRCGIH
13932	44300	A	14014	1	376	MQAVLLTVLQKHNQSVIYWY MKYIYYEGLVIMDDYDYVIMD YMTMETEKSHDLQAGGPGKPV VQFQFKPKILQASRGNGVKSWS KSKGLRTRCANVQGQENREGP AKHSPFLCLFVLFGSSTGRMIPN HTGDTSKVIAGTEERGVPKKVT LKFDACAAIDRPWAHQLDCTY HFKGSNDEHDSRLHTNAG*L*L C/RLWIT*LWRLRNPMICRLEA QESQWYSFSSSPKSCKPVEAMA LSPGPSPKA*EPGVLMSKGRKI GRAQLSIHLSSAFLFFSAPQQVE
13933	44301	Α	14015	1638	2289	VEKKKKPMSAPALRLPDLTKLF TLYVSETKKMAVRVLTQTVGP WPRPVAL\LSKQLDEVSKRWPP CPKSLVAIALLAQEADKLTLRQ NLNIKSPCAVVILINTKGHH*LM NARLARYQSLLCEHPRITLGSL QHLNPATFLPVSESPVKHNCVE VLDSVYSYVGPNHRDHPLNISR LGAVTWMGSSFHQPLQSDTLK KTTSPGSSHTRKLTGPRTAEA
13934	44302	A	14016		675	MAPPLRPLARLRPPGMLLRALL LLLLSPLPGLREGIGELITPIGT SLPDLDPARRRWEGGIGRVGSE VADLCPGKEGGKVPEAEKEGV WCFSELSFVKEPQDVTVTRKDP VVLDCQAHGEVPIKVTWLKNG AKMSENKRIEVLSNGSLYISEV EGRRGEQSDEGFYQCLAMNK\F*AILNQKAHLALSRIGST*RRRP DRP*EDEAFVMTTHCFQDLLTS
13935	44303	С	14017	18	329	
13936	44304	В	14018	391	1110	WASSDTPTGFTQRWRPDWAGG GGGEERGDGTGVVWEGEKVY RFRRVTGGSSFPSPAVLVRAPN TGMPKLTYIQT*GPLGERGKEP VPPN\PTLPQTFGGLLNASVVRK TEIPSPSPPPVCYLRASAERQ/PA GGSPQAKWRRAASSPPPLPPVH LPPPPPPPRPPPPPPP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13938	44306	A	14020	282	710	YPEPQGWRVAHTVQYPIHIQPL KPGPRGR*SIMDIQVPPKEGGV LILKN\SSRLLNSWTQGCLQISA FSPTSSTLPGTPWSNVGPYKIAP ITSFGE\PFIRVIWKQLPSWISIIQ V/SHKQ*QIHRSVDIGWSPGQEC GPKGCFV
13939	44307	A	14021		1416	MIILIDAEKAFDKIQQPFMLKTL SKLGTDGTYLKIIRAIYDKPTAN IILNGQKLEAFPLKTGTRQGCPL SPLLFNIGLEDLARAIRQEKEIK GIQLGKEEVKLSLFADDMIVYL ENPIVSAQNLLKLISNFSKVSGY KINVQKSQAFLYTNNRQTESQI MSELPFTIASKRIKYLGIQLTRD VKDLFKENYKPLLKEIKEDTNK WKNIPCSWVGRINIVKMAILP\R DIDQW/NRTEPSEIMPHTYNYLI FDKPEKNKQWGKDSLFHKWC WENWLAVCRKLKLDPFLTPYT KINSRWIKDLNIRPKTIKTLEEN LGITIQDIGVGKDFMSKAPKAM ATKAKIDKWDLIKLKSFCTAKE TTIRVNRQPTTWEKIFATYSSD KGLISGIYNELKQIYKKKTNNPI KK WAKDMNRHFSKEDIHAAK KHMKKCSSSLAIREMDIKTTMR YHLTPVRMAIIKKSGNNRCWR GCGEIGTL
13940	44308	A	14022	<u> </u>	3450	GCGEIGTE
13940	44309	В	14022	1	3171	
13741	77309	ما	17023	l*	121/1	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	•	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13942	44310	A	14024		2433	MRTKTQHTRISGTHSKPVCRGK FIALNAHKRKQEKSKIDTLTSQ LKELEKQEQTHSKASRRQEITKI RAELKEIDTQKTLQKINESRSW FFERINKIDRPLARLIKKKREKN QTDTIKNCKGDITTDPTEIQTTI REYYKHLYANKLENLEEMDKF LNTYTLPRLNQEEVESLNRPITG AEIVAIISSLPTKESRTGWIHSRI LPEVQGGTEKEGILPNSFYEASII LIPKPGRDATKKENFRPISLMNI DAKILNKILAKRIQQHIKKLIHH DQVGFIPGMQGWFNIRKSINVI QHINRTKDKNHMIISIDAEKAF DKIQQRFLLKTLNKLGIDGTYF KIIRAIYDKPTANIILNGQKLEAF PLKTGTRQGCPLSPLLFNIVLEV LARAIRQEKEIKGIQLGKEEVKL SLFADDMIVYLENPIVSAQNLL KLISNFSKVSGYKINVQKSQAF LYTNNTQTESQIMSELPFTIASK RIKYLGIQLTRDVKDLFKENYK PLLKEIKDDTNKWKNIPCSWVG RINIVKMAILPKNWKKTTLKFI WNQKRAHITKSILSQKNKAGGI TLPDFKLYYKATVTKTAWYW YQNRDIDQWNRTEPSEIMLHTY NYLIFDKPEKNKQWGKDSLFN KWCWENWLAICRKLKLDPFLT PYTKINSKWIKDLNVRPKTIKTL EENLGITIQDIGMGKDFMSKTP KAMATKAKIDKWDLIKLKSFC
13943	44311	A	14025		1023	MGAIYDKPTANIILNGQKLEAF PLKTGIRQGCPLSLLFNIVLEV LARAIRQEKEIKVIQVGKEEVK LSLFADDMIVYLEDPIISAPNLL KLISNFSKVSGYKINVQKSQAF LYTNNRQTESQIMSELPFTIASK RIKYLGIQLTRDVKDLFKENYK PLLKE\KEDTNKWKNIPCSWVG RINIVKMAILP/KAICRKLKLDPF LTPYTKINSRWIKDLNVRPKTIK TLEENLGNTIQDIGMDKDFTSK TPKAMATKAKIDKWDIIKLKSF CTAKETTIRVNRQPTEWEKIFAI YSSDKGLISRIYNELKQIYKKKS NNPIKKWAKDM/NRHFSKEDIY AAKRHM
13944	44312	В	14026	1	2206	
13945	44313	Α	14027	1	1689	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13946	44314	Α	14028	1	2523	MKQFLLYLDESNALGKKFIIQDI
į						DDTHVFVIAELVNVLQERCHTR
		ŀ				LGYTEFLVAWRVTFGLCVEAV
1						TLHLKYQILIRGLLEMMSFSDA
1						DILKQLPVTVPGLFPASLSPSSL
						LGNSPPSWLRHNSESKVSAVSS
						PSATKTLSTGIGKLDPGHKEMA
						EESELLKNKMQAPPLSRCPESQ
						KCQHQLRLHHWKPSVRHQVKR
		ŀ				RSPAVLRSAMPPADCPAVLEAT
i						TATHPEKGTALSKHLPSSDSMS
						LKVDVEALENSPGATYIWKGG
						KVTRDSQPKEQGKGDLKKKKK
						GKLPKNYDPKLTPDPERWLPM
1						QECSFYQGRKKGKKKDQMGK
ŀ						GTQGATAGASSELDARKTVSSP
						PTSPRPGSAATLSASTSNIIPPRH
						QRPAGAPATKKKQQQKKKKG
						GKGFPVLREITVVKVDTLVVFQ
						ILEERLSVFHIQYDTSYPFSTVDI
ľ						EDHECAVWLLLRKSKSDDKTT
	:					RLEAVREMSETHHWHDAEKAF
						DKIQQPFMLKTLNKFGVDGTY
						LKIIRAIYDKPTANIILNGQKLE
						AFPLKTGTRQGCPLSPLLFNTV
						LEVLARAIRQEKEIKGIQLGKEE
						VKLSLFAGDIIVYIENSIVSAPKL
						LKLISNFSKVSEYKINVQKSQAF
,						LYTNNRHTESQIMSKLPFTIATK
						RIKYLGIQLTRDVKDLFKENYK
						PLLNEIKEDTNEWKNIPCSWVG
						RINIMKMAILPKVIYRFNAISIKL
13947	44315	A	14029	1	2868	
13948	44316	A	14030	1	3099	MGELITPLSTLDRSTRQKVNKD
						TQELNSALHQGDLIDIYRTLHP
						KSTEYTFFSAPHHTYSKIDHILG
1						SKALLSKCKRTEIITNYLSDHSA
						IKLELRIKNLTQNRSTTWKLNN
						LLLNDYWIHNEMKAEIKMFFET
						NENKDTTYQNLWDAFKAVCR
						GKFIALNAHKRKQERSKIDTLT
						SQLKELEKQEQTHSKASRRQEI
						TKIRAELKEIETQKTLQKINESR
						SWFFERINKIDRPLARLIKKKRE
						KNQIDTIKNDK
13949	44317	Α	14031	2	3419	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	I.	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
l				sequence		
13950	44318	A	14032	1	2685	MGDFNTPLSTLDRSTRQKVNK
						DTQELNSAPHQADLIDIYRTLH
-						PKSTEYTFFSAPHHTYSKTDHIL
						GSKALLSECKRTEHTNYLSDDS
				:		AIKLELRIKNLTQNRSTTWKLN
						NLLLDDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFVALNAHKRKQGRSKIDT
				-		LTSQLKELEKQEQTHSKASRRQ
					1	EITKIRAELKEIETQKTVQKINES
						RSWFFERINKIDRQLARLIKKKR
}						EKNLIDAIKNDKGDITTDPTEIQ
1						TTIREYYKHLYANKLENLEEM
		}				DKFLDTYTLPRLNQEEVESLNR
						PITGSEIVAIINSLTTKKSPGPDG
						FTAEFYQRAIRQEKEIKGIQLGK
ľ		ŀ				EEVKLSLFADDMIVYLENPIVS
						AQKLISNFSKVSGYKINVQKSQ
						AFLYTNNRQTESQIMSELPFTIA
		:				SKRIKYLGIQLTRDVKDLFKEN
						YKPLLKEIKEDTNKWKNIPCSW
		:				VGRINIVKMAILPKVIYRFNAIPI
						KLPMTFFTELKKTTLNFIWNQK
						\RAHIAKS/VLSQKNKAGGITLP
						DFKLYYKATVTKTAWYWYQN
						RDTDQWNRTEPSEIMPRIYNYL
						IFDKPEKNKQWGKDSLFNKWC
						WKNWLAICRKLKLDPFLTPYT
						KINSRWIKDLNIRPKTIKTLEEN
		İ				LGITIQDIGMGKDFMSKTPKAM
			į			ATKAKIDKWDLIKLKSFCTAKE
						TTNRVNRQPTKWEKIFATYSSD
13951	44319	Α	14033	845	1616	ARAEVKLSLFADDMIVYLENPII
						*ARAEVKLSLFADDMIVYLENP
						IISAQNLLKLISKFSKVSRYKINV
						QKSQAFLYTNNRQTESQIMSEL
		ł				PFTIATKRIKYLGIQLTRDVKDL
						FKENYKPLLNEIKEDTNKWKNI
						PCSWIGRINIVKMAILPKVIYRF
						SAIPIKLPMTFFTELEKKNWLAI
						CRKLKLDFFFIPYTKINSRWIKD
						LNVRPKTMKTLEESLGNTIQDI
						GIGKDFMTKTPKAMATKA/KKS
						FCTAKETTIRVNRQPTEWEKIF
	<u> </u>	<u> </u>				AIYPSDKGLIS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13952	44320	Α	14034	707	2878	TLMQKSSIKYWQNESSSTSKSL
						STMIKWASSLGCKAWFNIRKSI
						KVIQHINRAKDKNHMIISIDAEK
	ļ					AFDKIQQPFMLKTLNKLGIDGT
						YFKIIRAIYDKPTANIILNGQKLE
						AFPLKTGTRQGCPLSPLLFNIVL
						EVLARAIRQEKEIKGIQLGKEEV
						KLSLFADDMIVYLENPIVSAQN
						LLKLIRNFSKVSGYKINVQES\Q
						AFLYTINRQTESQIMSALPLTIA
						SKRIKYLGIQLTRDVKDLFKEN
		ĺ				YKP\LLKEIKEDTNK\WKNIPCS
		l			ŀ	G\EGRINIVKMAILPKNWKKTTF
İ		1				\RFIWNHKRACIAKTILSQKNKA
		1				GGITLPDFKLYYKATVTKTAW
						YWYQNIDIDQWNRTEPSKIIPRI
	1					YNNLIFDKPDKNKKWGKYSLF
						NKWCLENWLAICRKLKLDPFL
						TSYTKINSRWIKDLNVRPKTIKT
						LEGNLGNTIQDIGMGKDFMSKT
						PKAMATKAKIDKWDLNNLKSF
						CTAKETTIRVNMQPTEWEKIFA
						IYPSDKGLISRIYNELKQIYKKK
						TT\NPIKKWAKDMNRHFSKEDI
						YAAKKHMKKCSSSLAIREMQI
						KTTMRYHLTPVRMAIIKKSGNN
						RCWRGCGEIGTLLHCWLDCKL
						VQPLWKSVW*FLRNLELEIPFD
						PAIPLLGIYPNDYKSCCYKDTCT
			1			RMFIAALFTIAKTWNQPKCPTII
			}			DWIKKMWHIYTMEYYAAIKND
						EFVSFVGTWMKLEIIILSKLSQE
13953	44321	Α	14035	869	3869	RHKKPFKKLTNPGAEIQTTIREY
				·		YKHLYANKLENLEEMDKFLNT
						YTLPRLNQEEVESLNRTITGSEI
1						EARINSLPTKKSPGPDGFTAEFY
						QRYKEEMVPFLLKLFQSIEKEGI
						LPNSFYEASIILIPKPGRDTTKKE
						NFRPISLMNIDAKILNKILANQI
				ĺ		QQHIKNLIHHDQVGFIPGMQG
1				1		WFNICKSVNVIQHINRTKDKNH
1						MIFSIDAEKAFDKFQQLFMLKT
1		1				LNKLGIDGMYLKIIRAIYDKLT
1					•	ANIILNGQ
13954	44322	Α	14036	2	99	
		<u></u>	1	I	1	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	Ì	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13955	44323	A	14037	1	914	TVTVIHTVARVGNLDIQIQRKS
						ELFLSLVVCEAIKNEMNVMLKS
						SPVDLVTATDQKVEKMLISSIK
						EKYPSHSFIGEESVAAGEKSILT
				ĺ		DNPTWIIDPIDGTTNFVHRFPFV
				İ		AVSIGFAVNKKKKVDETYLGL
						VKETCFPLQIEFGVVYSCVEGK
				ŀ		MYTARKGKGAFCNGQKLQVS
				ĺ		QQEDITKSLLVTELGSSRTPETV
						RMVLSNMEKLFCIPVHG\IRSVG
				İ		TAAVNMCLVATG\GA\DAYYE
						MGIHCW\DVAGAGIIVTEAGGV
						LMDVTGGPFDLMSRRVIAANN
						RILAERIAKEIQVIPLQRDDED
13956	44324	В	14038	91	1959	
13957	44325	Α	14039	28	152	LPSPSSKDTDS/AASCCAPRGRA
		$oldsymbol{ol}}}}}}}}}}}}}}}}}$				KGSVSNSRLSFPPSSVPLP
13958	44326	Α	14040	35	879	GSQCSSPPPASRCGSCSSAPSRH
						CPCRDFEPAVAGLARTPVLTTK
						TNPSPPARTGLNAPSMGTSLI/P
						ALCCFPLYQGSTELQCKAP**LP
						SPSSKDTDS/AASCCAPYFLVLL
	1					KVLSSVDNSSIWCSAGPPAQS/S
						SLYTIPMSSMSAKNCSVLRSGT
		ı				WASFTCS*GSQRVSLGTPEALG
						TEDHTSRSLGHSHVCHVDLSPA
						TASCSFQTASSEVSPPATASCSF
						QTASSEVSPPATASCSFQTASSE
						VSPAATASCSFTFSFFFVCALWF
12050	144207	 	14041		159	GICPSCSAAVLST EHQCENPQ*NTGTPNPAAHQK
13959	44327	A	14041	3	139	AYPP*SSGLHPWDNQRQKPHD
	İ					YLNRCRKGL
12060	44220	 	14042	54	245	I
13960	44328	Α	14042	34	245	GGGGGEAEDRLREPETEKALSS SL/RRAPDQC/RALIMQLFQAHC
						FFLST/QATAALQAHYAHIFPSK
13961	44329	A	14043	903	1205	WDVSQNNKSYL*QTHSQYHTE
13901	44329	^	14043	303	1203	WAKTGSIPFENWHKTGMPSLT
						TPIQHSVGSSGQGNQTNREPNH
						E*TPIHNCFKENKIPRNPTYKGC
						EGPLQGELQTTAQ
13962	44330	В	14044	1	339	LOI LOODLOI IAO
13962	44331	В	14044	1	1547	
13703	144331	Б	14043	<u> L'</u>	1347	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
13964	44332	A	14046	807	1440	I INTGKPNPAAHRKAYPPRSSRLH
1570 1	111332	`	1 10 10			P*DLQQTPTDLQLSDLTPGRKT
						NKQKGIASPSTKRTSTPKPHLQ
						ANAQRFCHHQACVTRVPKGST
						KHGKEQLVPATAKTCQIVKNIN
						AMKKLHQLTEIQITIREYHKHL
						YANKLKNLEEMDKFLDTYTLP
						RLKQEEVESLNRPIMGSEIEAIIN
						SLPTKKSPGPDGFTAKFYERYK
]						EELVSFLLKLFQSIEKEGILPNSF
						YEASIILIPKPGRDTTKKENFRPI
						SLKNIDAKILNKILANRIROPIEK
						LIHHDQVGFIPEMKGWFKKCKS
						INVIHHINQTNDKNHMIISIDAE
						KAFDKTQQPFMLKTLNKLGTIR
						QKKEIKGIQLGKEEVKLSLFAD
						DMIVYLENSIVSAQNLLKLISNF
				İ		SKVSRYKINLQKSQAFLYTNNR
						HMESQITSELPFTIDTKRIKYLGI
						QFTRDVKDLFKENYKLLLNEIK
						EDTNKWKNIPCSWIEESIS
13965	44333	Α	14047	603	764	LPLFLIEYPLFLSLA*LPWPKLPT
						LC*IGVVREGIPVLCCFSKGMLP
						VFAHSV
13966	44334	Α	14048	1566	1865	MQSIRKTIGIHAKYDKACRNGE
						DICLRETNSAADTQAKRV*SGP
						PANSNRPAAEGPDC*KEN*QTE
						RTSTPKPHLYVTIIKDQRGTQLL
						TNNGTKLDRE
13967	44335	В	14049	1	1932	
13968	44336	Α	14050	711	1575	TRKFRRNG*IP*HIHPPKTKSGR
				1		SRI\FNRPITGSEIEAIINSLPTKKS
			ļ			PGPDGFTAEFYQRYKEELTNIL
						DEHRCKILNKILANGIQHDIKKL
		İ				IHDDQVGFIPGMQGWFNIRKSI
						NVIQHVNRTRDKNHMIISIDAE
				Ţ		KAFDKIQQPFMLKTLNKLGIDG
						MYLKIIRAMYDKPTANIILNGQ
		1				RLEAFPLKTGTRQGCPLSPLLFN
						IVLEVLARAIGQEKEIKGIQLGK
						EEVKLSLSADDMIVYLEKPIISA
						QNLFKLISNFSKVSDTKSMYKN
						HKHSYTPITDKQRAKS

SEQ ID	_	1	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	Sequence			sequence		,
13969	44337	I A	14051	<u> </u>	1579	 MQGWFNVRKSLNIIHHINRNND
13909	44337	A	14031	1	1379	1 7
				1		KNHIIISIDAEKAFDKIQQPFML
						KTLNELGIDGTYLNIIRTIYDRP
						AANIILNVQKLEAFPLKTGTRQ
						GCPLLPLLFNIMLEVLARAIRQE
						KEMKGIQLGKEEIKFSLFADDII
				-		VYLENPIFSAPNFLKLISNLSKFS
				İ		GYKINVQKSQAFLYIINRQTESQ
						IMSELRFTIATKRIKYLGIHLTR
	1					DVKDLFNENYKPLLNEIKEDTN
						KWKNIPCSWTGSINIVKTAILPK
						VIYRFNAIPIKLQLTFFTELEKTA
						LKFMWIQGYSNQNSMVLTLLV IPRKMRSAVELQQTSTDLQLRD
		•				LTVRKKTNKRKATASTSQKGH
						LHQNPICRSPTSKTKEVGFRRSV
						ITNFSELKEDVRTYHKEAKNLE
						NRLDEWLTRINSVEKTLNDLKE
						LKTMARDLHDACTSFNRRFDQ
						VEERVTVTEDQINEINDGENGT
						KLENTPQDIIQENFPNLA\SRPTF
						KFRKYREHYKDTPREEQPQDT*
						LSDSPRLK*RKKC*RQPERKVE
						LPTKGSPSD
13970	44338	Α	14052	50	1105	TRKSRRNG*IPGHIYPPKTK\QE
		1				EGESLNRPITGSEIEAIINNLPTK
						KSPGPDGFTAKFYQSTNNKNH
						MIISIDAEKAFHKIQQPFMLKTL
						NKLGIDGTYLKIIRAIYDKPTAN
						IILNGQKLEAFPLKTGTRQGCPL
						SPLPFNIVLEVLARAIRQEKEIN
						CIQLHNEEVKLSPFADDMIAYL
		1				ENPIVSAQNLLKLISNFSKVSGY
		ļ				KINVQKSQAFLYTINRQTESQI
		ŀ				MSELPFTIATKRIKYLGIQLTRD
						VKYLFKENYKPLLNEIKEDTNK
ŀ						WKYIPCSWIGRINIVKMAILPKV
ļ						IYRFNAIPIKLPMTLSSQNWKKT
					İ	TLKFIWNQKRARIAKSILSKKN
		<u> </u>		<u> </u>		KAGSIMLPDFKL
13971	44339	Α	14053	1	764	MQGWFNVRKSLNIIHHINRNND
						KNHIIISIDAEKAFDKIQQPFML
		1				KTLNELGIDGTYLNIIRTIYDRP
					[AANIILNVQKLEAFPLKTGTRQ
						GCPLLPLLFNIMLEVLARAIRQE
						KEMKGIQLGKEEIKFSLFADDII
				1	[VYLENPIFSAPNFLKLISNLSKFS
					}	GYKI*FPLFLIEYPSFLSLA*LPW
	1	1	I	<u></u>	l	PALPTLC

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13972	44340	A	14054		344	MGINQRRKAENSKNQSASSPTK DCSSSPAKEQSWTENDFEELTE VGFRRQANAERFCHHQVCPKR APEESTKHGKKQ/RSISHCKNM PNCKDHRCYEETASTDSLAETQ QQKREF
13973	44341	A	14055	301	370	DDGVF*IDNHVICKQRQFDFLFS
13974	44342	A	14056		1418	MVREGFPEEVTFELKNELEISK GQQMGKNIPDTGASSAKALGE RELGTLDKMKNSPSDGSKKDIR THRKEAKNLEKRLDEWLTRINS VEKTLNDLMQLKTMARELHDA CTSFNSQFDQVEERINKIDVRLA RLIKKKREKNQIDAIKNDKGDIS TDPTEIQTTIREYYKHLYANKL ENLEEMDKFLDTYTLPRLNQEE FESLNRPIIGSEIEAVSNSLPTKK SPGPDGFTAEFYQRYKEELVPF LLKLFQSIEKEGILPNAFYEASII LIPKPGRDTTKKENFRPVSLMNI DAKILNKILANRIQQHIKKFIHH DQVSFIPRMQGWFNICKSINIIH HINRTNDKNHMIISIDAEKAFDR IQQPFMLNTLNKLGIDGMYLKII RTIYDKPTANIILNGQKLEAFPL KTGTRQGCPLS/ATPVQHSVGS SDQGSQARERNKGYSIRK*GSQ IVSVCR*HDCIFRKPHRLSPKSP
13975	44343	A	14057	1240	1596	CYSFLFVRCPSDSQVPQLQVC/C EFAGGPLQTVFAWVSPAEVVE MFLQG*RKPFYMSY/TSFEMAL MKVWTMVRTSASPVGDVFGR GWSPFPTSTVGALTVFGVSPGS CRSSPLPSERLWV
13976	44344	A	14058	507	971	PANQKKSRTRRIHSQILPEAQR GAGTIPSETIPINGKRGNPPKLIL *SQHHPDTKAWQRHNRKREFY TDIPDDH*CKNPQ*NTGKPNPA AHQKAYPPRSSWLHLWDARLV QRTQISKRNSSYKQNQRQKPHD YLSRCRKGLQQNSAALHAKNS
13977	44345	Α	14059	561	758	
13978	44346	A	14060	2364	2663	DGQLTLVSILNHYSLAQSCEY* DGECFCLHCSASPFL*EGAQST ACTAPPECFASDLGAV*LPQHS WCSTLLLEGAPWPGTLGTAPQF QHRKLGTNLAG

SEQ ID	SEQ ID NO:		SEQ ID NO:	1		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	or peptide sequence	deterion, \-possible nucleotide insertion)
13979	44347	Α	14061	1	64	MPIGKVGEYYIKGTLRATKESE
						QQSSALELSSDRVYLNEKEPGD
						QPWNAAFVFRGCGVALALRPP
						HLGWGEPRWAGFRGAFGFPFS
						TLNPSNWSLTLIIQGCKQEGFYS
1		ĺ				PSMRPRVVPKRGSAPKVRAEEN
					:	AALPSRCPPGPLPVAQP/TGPAD
1						GGSCEQCPSCGCGCCQLPPPLP
		ĺ				WFPR*TPCDKRV
13980	44348	Α	14062	1	774	MFVWSNVEGHSVAMFPWYSIP
						FLNPPCSHTRPSNLPVTQWPPTR
						ENNLPSWQLLLTSIHQAQFLSA
						LHKEQGSSSEKDGRSPNKWDK
						DHIRCPMS\AVMIFSKRHQAL/V
1						RAHQGHPNQDNWTVSQMLSK
		ı				WWYTLGPNERQKYHELAFQAT
						AALQAHYVHIFPSKGRWELLH
						GSQRLDWGRCQLKLLGQQGQ
						DVLHTNLGKEENSSCWGYKME
						CEEGFGPQGGDCGLGGVHAPL
						LSPEGLDSEQNTLIMGDIFHCA
						KKLP
13981	44349	В	14063	1	753	VDDDDDDD ACYU ACDAYU ACDA
13982	44350	A	14064	2	1457	YPRRRPSRAGVLAGPAVLAGPP
					-	SSPGRRPRRAAFFAEPPFSARRS
					+	PRRAAVLAVPPSSPRRPRRAA
						VLAAPRVLAAPPSSPRRRPR\PP
						PSSPRRPRRAAAGSLPGAELP
						QSSRLAAHL\AAD*PDS*IAAGA VPRGWEGLTIMAEGKEEQVTF
						•
						YVDGSRQRAYAEKPPVVKTIRS CTV/SGHSRHNSSSGM*/SSPNSP
						/IPFQFTFQIPSLLQAALWLCL*V
		İ	ļ			TWHFFVR*PVLSSRHCFSPGGLF
				:		SLSLQSAGCPAGSLQAEFVALP
						PGGQSRLHRTRAS*VPGAFQEK
		ł				CPLERVWPFHPSKQHPRNGLAF
						PLP*APQRTQPAEDTLPRHPEM
		l				GLILSQGTAG/PGRECQATQLPA
		l				QHSHAEVLHFGGAMSGQLSLV
						GPQDSKRTARLTDSQHLGSPT*
		1				OPPGLTEGTSLEOG**S*MGWM
						VGVIAGAPSGWSGSSGPSEGTP
		1				LFSSPA*LPWPELPTLC*IGVVR
]				EGIPVLCQFSKGMLPVFAHSA
13983	44351	В	14065	72	1235	Zer (Degrending) (Tritish
13984	44352	В	14066	1	2715	
		1		L		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
13985	44353	Α	14067	50	1304	TRKFRRNG*IP*HIHPPKTKSGR
		· ·		,		SRI\FNRPITGSEIEAIINSLPTKKS
						PGPDGFTAEFYQRYKEELTNIL
						DEHRCKILNKILANGIQHDIKKL
1						IHDDQVGFIPGMQGWFNIRKSI
						NVIQHVNRTRDKNHMIISIDAE
						KAFDKIQQPFMLKTLNKLGIDG
ŀ						MYLKIIRAMYDKPTANIILNGQ
						RLEAFPLKTGTRQGCPLSPLLFN
						IVLEVLARAIGQEKEIKGIQLGK
		l				EEVKLSLSADDMIVYLEKPIISA
						QNLFKLISNFSKLSGYKINVQKS
						QAFLYTNNRQTESQIMSELPFTI
1						ATKRVKYLGIQLTRDVKDLFKE
			į			NYKPLLNEIKEDTNKWKNIPCS
	,	1				WIGRINIVKMAILPKCKTVGLLS
						NISTHKSAVHALVEAGGIPSLIN
						LLVCDEPEVHSRCAVILYDIAQ
1						CENKDVIAKY
13986	44354	Α	14068	1	1155	MFVWRNVEGHSVAVFPWYSIP
13980	44334	^	14000	1	1133	FLTPPCSHMRPSKLPVTQWPPT
						RENNLPSWQLLLMSVHQAQSL
						SALRKEQDSSSEKDGRSPNKW
						DKDHIWWPMS\AVIIFSKQHQA
						L/VRAHQGHPNQDNRTISQMLS
						EQWYTLGPNEMQKYDLAFQN
						VEARIRY/CHRPRKRARFLSHAS
						PPWRQLPAAGAEASGAWNLPA
						GGG*HDYNRCSVPVFRTEVSTR
						PRPSEICGLTQE*TLLSAKSQVG
						NTFGFAGHTISDTTTPCGLCTSK
1		İ				AAIEVPEADTLPHICSWHKGEE
						CGLRGPEPSSPPPRPSLAARVSA
						GRGALRRRRRPEVPP\SRAPRSA
1						ATATPANEQRAVADVGQWAR
						ARGGRVRAAGRSGLLRVKMM
1222	11055		14066	1221	1504	KMKTVMIHFHLMKDEYIFSS
13987	44355	Α	14069	1331	1584	SLQRKAARFASSSTVPRRPQGIL
						PIISFSNVPVGSRRLKAPLVFIGP
						G*ITLNRIPYLPHSAARDPFTWN
			1.40=6		1040	TFSFLLPCEEVPST
13988	44356	В	14070	1	1248	
13989	44357	В	14071	1	459	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13990	44358	A	14072	1	2156	MGKKQSRKTGNSKNQSASPPP KERSSSPAMEQSWMENDFDDL REEGFRQSNYSELKEEVRTHGK EVKNLEKKLDEWLTRITNAEKS LKDLMELKTTARELHDECTSLK RVSVTEDQMNEMKREEKFREK RIKRNEQSLQEIWDYVKRPNLH LIGVPESDRENGTKLENTLQDII QENFPNLVRQANIQIQEIQRTPQ RYSSRRATPRHIIVRFTKVEMK EKMLRAAREKGRVTHKGKPIR LTADLLAETLQARREWGPIFNI LKEKNFQPRISYPAKLSFIDRST RQKVNKDTQELNSALHQADLI DIYRTLHPKSTEYTFFSAPHHTY SKIDHIVGSKALLSKCKRTEIIT NCLSDHSAIKLELRIKKLTRNHS TTWKLNNLLLNDY WVHNEMK AEIKMFFETNENKDKTYQNLW DTFKTVCRGKFIALNAHKRKQE RSKIDTLTSQLKGLEKQEQTHS KASRRQEITKIRAELKEIETQKT LQKINESRSWFFEKINKIDRPLA RLIKNKREENQIDAIKNDKGDIT TDPTEIQTAISEYYKHLYANKL
						ENLEEMDKFLDTYTLPRLNQEE VESLNRPITGSEIEAIINSLPTKK SPGPDGFM/RRILPEGQGGAGTI PSETVPINRKRGNPP*/PHFM/SA SIMLIPKPGRDTTKKENFRPISL MNIDAKI/RQ*NTGKPNPAAHQ KAYPP*SSGLHPWDARLVQLTQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
13991	44359	A	14073	3	1794	AMLPMELGCGPLPEPLPVGCSR FSLFK*QTCISTVP/GYMVTAQS MSSTPPPPSPSTLPSSPSPPPPPLPQ PLPPPPPSPPTLSSLSSPSPPRPPL VSPSTLPSPQPSSPQPLLPPSSSPP SLPSPPPPSPPLPSPSPSAIPSLPPP SPQPLPPPPPSSPPPSLPSPLPPP PLSSSPSSPLSPSPPPSPP
						DGRSPNKWDKDHIWWPMSGG HDLQQAAPGPGRAHQGHPYQD NWTISQILSERWYTLGPNEMQK YHDLAFQHMAGEDIASDEEHM VIHEEEGVMVSLLMTALAPLTL ISSSRIFGKVYGPTPSSSYTYSD ASSSTLAPTSFLLGPGAFKAQES GEEAEDGLRELETEKALSSSL/R RALDQ/*LALIMQLFQAHCFFLS T
13992	44360	A	14074	4217	2496	MLLSIDAEKAFDKIQQPFMLKT LNKLGHGYEQHHLHTTNDVDE EDLSDAASKGDDFALSEQSQD AHFLQPEAYGLGEGAETATGT AHQGNHVRVEECGRSLCGCVP LVLHPLPDPSLQPHEAQQPASH SVACNQRKQPAKLPAVAHERP PGGTGSVDPGRPPGATCPESPG PATPHTLGVVEPGKSSPPTMEE EPWAPQGSPCWTAQSLSALRK EQDSSSEKDGRSPNKWDKDHI WWPMS\ALMIFSKRHQAL/V
13993	44361	A	14075	4317	5099	KMFVFLCISAQLSLQCFRGSFHF IKIYDFSFQLFFLFLELPYLLLN HFKMLELVLQQGHPNQDNWT VSQMLSKWWYTLGPNERQKY HELAFQVKVAH\CNKD*KKFSS EAKPTSQGLAG\GNKGSWEWS MSETGTATAPGVSSELLSVAAQ TLQSSDTK\SSFCGAEWGHP*GI GCDDVIADDGFSTTDTDLKFKE WVTD\ESGDNSGEEPEGNKGFG GKVFAPVIPSSFT/HCRPLLDPE/ PPGSPDPPAAFGKVYGPTLSSSY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
12004	144262	<u> </u>	14076	 	637	I MSRVGSCRWVCGLADFKNGAT
13994	44362	A	14070	1	037	ł
						DLCDKSLHNRELSAERPLNEQI
						AEAEEDKIKKTYPPENKPGQSN
						YSFVDNLNLLKAITEKEKIEKER
						QSIRSSPLDNKLNV/EKDVDFNQ
						EFGKLIDGY\DSTKEWDWVHKF
						QDIPNVLVGTAFNVDDGNGGF
						VVHWLNNKEFHFTSSTEVFMH
						QLRKLSDKQVDHENDDADRED
		ļ				EEHSQEDRERGLHMKLDH
13995	44363	Α	14077	448	464	KKRA*GSVSQRFPAET*LAD*VI
		<u> </u>				NTGSQEAKRREQR
13996	44364	Α	14078	179	382	NTPNQRSVASNRFVKCTHQHS
	ļ					VKCTQSALCKMHQSAGFLKVD
		1				NHRED*KKGHSDRAKTEHGRG
	_					QIRE
13997	44365	Α	14079	697	1654	PKPSVGRMGFLGTGTW\ILVLV
						LPIQAFPKPG\GSQ\DKSL\H**ED
					:	LSAERPFELNRLLEAEGRTRFK
						KTYPP\ENKPG\QSNYSFV\DNL
						NLL\RAITEKEKIEKERQSIRSSP
						LDNKLNVEDVDSTKNRKLIDD
						YDSTKSGLDHKFQENETADNSF
1						SQEEEPVVAGEDLPPPSPQESD
						VQPVQPEEVSARDFLQRLDLQI
						KLSTQAAKKLKEESSSLERMRE
						ETSLSPTVPTKRWDTRILLIGAF
		l				YRVLIGAFYRVLMGAFYKPIAS
						YRALIGVFYRALMGAFYNPLV
						RQKSSPRPHSTQEVQLASPLTN
						TQITYYVIFTV
13998	44366	Α	14080	72	1117	ILENHLSDRVAEGGPLQGLRLG
		1				VQGEAVPPGLVHQRGGLGPLD
						GELIGSVGQQVSDVEGAGSHR
		1				G*RHTGRTPAVSAGPAA*SATH
						TAPMWLAGW\YEPAQPGPVLV
						SPPTPFGQTRQASWCSA*ARCA
	1					GSWWH*HQTPGGA*RVAAPAP
						SWSRSCP*RPTRSPAAAPPAPSR
						PGRAVGLATGAGHPAQHGHGR
						AVPTGPRAAGAGSWSELCPPSS
						CAGTSRGR*AGTRR*AAPAGSG
		1				RPGPGAWAGPRCSRSTYAAAA
						CGRGGA*AGADSGPRSGTHAG
						HCGARCRSSTPLVPAAAGHLPP
						PGPGRTGDCPASSCPGPGLPHG
						CSLSGSWGEGWVSAARQAGLP
						WAVA
13999	44367	A	14081	2	269	
14000	44368	A	14082	1	591	
14001	44369	A	14083	2	477	
1.1001	1,,,,,,,	1,,	1. 1003	1-		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14002	44370	В	14084	237	419	
14002	44370	A A	14084	2	1220	 LPPGFVMAAAAARWNHVWVG
14003	44371	^	14003	2	1220	TETGILKGVNLQRKQAANFTA
		1				GGQPRREEAVSALCWGTGGET
						QMLVGCADRTVKHFSTEDGIF
						QGSETLPG\GEGMFRGLAQTDG
		ľ				TLITCVDSGILRVWHDKDKDTS
İ						F*PTPGTESGPWGCVGCAKTQA
						HPHVVCHRLGKRMLLKIWDLQ
						GS\GNLCFKAKNVRNDWLDLR
						VPIWDQDIQFLPGSQKLVTCTG
						YHQVRVYDP\ASPQRRPVLETT
						YEEYPLTAMTLTPGGNSVIVGN
						THGQLAEIDLRQGRLLGCLKGL
			}			AGSVRGVAV\HPSKPLT*PPVRL
						DRVLRIHRIQNPRGLEHKVYLK
						VSIE/RALLLSGRDNLGG*APEP
		1				SKTQTRCP*EDTEDR*TLGDPW
		-				RQLPSGKLFGVWSSPKELSKRD
						GERRSGLGPPAPDAPVPTL
14004	44372	В	14086	96	592	
14005	44373	Α	14087	102	612	HCAGSPHSPRSLSGTPLPGLPL
						WRHLRSPSAHRCTVGAPFWAG
	1					QDRSQLP*LAGRCGGRGPSGNP
						GCA\PACGPAGVPGGRGLGGPR
						TRSSQPALPAPGNEGLSTRASG
•		İ				CRGRTGSPSSASPPLGLSCLPAG
						QGWGP/GSPPCVSLPPTPWAPV
14006	44274	С	14088	1	636	RPEPPG*AAPPAPRRPVP
14006 14007	44374	В	14088	31	612	-
14007	44376	A	14090	234	504	SFYHLGAGE*PGKSRVPIPQFQS
11000	1,370	``	11000			PWGPYCQGRARGGEVCHLQAV
						PPRGAGAARTLSGHPLGSPEQL
		ŀ				PQPGPAAELPGGVVPRSTPPCP
14009	44377	В	14091	1	924	
14010	44378	Α	14092	11	354	RLSTSPPDSGAQLASPSGSHTRA
						AGGAACQSRAVRS/PFPSPWVV
						DGTGCGGAGGGARWGGSGCT
						GAHGVGGR\PGMAG*GPEPCP
						EGRQLRPLAPVGCGQ*REP\G*K
						NSMGETTT
14011	44379	A	14093	147	513	GSASGVVRLSRWARGLAGFRS
						EAADLRGECYSS*GSASGVVRL
						SRWARGLAGFRSEAADLRGEC
						YSS*KQRGPKELRSPAGFT*WIP
						/LPGLQVELPASPAPCAPTPQPL
						SG*WD\GRRGAGGGARRGGSG
1						RTGALGVG\GGSGMAGCRSRA
	<u> </u>					LPPREGS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14012	44380	A	14094	418	1026	WNPHQGCRWSCLPVPRHALAF
14012	14300	 	114024	710	1020	LSPWVIDGTGRHGAGGGALWG
						GSGRTGAHGVGG\S*GMAGCR
						SRALPHGKAAKAR*EIERSASG/
						TGTAGGPSTPSAATGPGAKSLI
						ARGQQGWPAAPSAGPAKPTPT
					i	GNSSWPASAAGSSGSCSRLS/SP
						HLPAS*GSGLQPWPAQKGAPTV
						QQQAEGLPIPRLPEEWISLPFLG
						HQGEHSGMA
14013	44381	A	14095	897	2111	PPLLQSSPNFFLICYLPWHNSHE
	11301	`	1 10,5	,		NTHRTGRRGAGGGARRGGSGR
						TGAHGVGGRLRHGGLQVPSPA
						PWEGS*GPARNRAQRRYHCSCP
		l				GLQSGLPHYSGYHT*PDEKEKN
						GREGQKLER
14014	44382	A	14096	1304	1922	KLNIGVIYSSEIPDSGAQLASPS
						GSGTRAAGGAVCQSCALRSHSS
						ALGWSMG/PGRRGAGGGARW
		1				GGLGRTGAHGVGGRLRHGGLQ
	:		Ì			VPSPAPREGS*GSVRNPAQRRW
		1				AGTAGGPSTPSAATGPGAKSLI
						TRGQQGWPAAPSEGPAKPTPTR
		1				NSSWPASAARSPGSRSR/RLPPH
						LPAS*GSGLQPWPAHYFEFFLPI
					1	GQNAQTEVSY
14015	44383	Α	14097	318	533	MVCFDGADCPWVVDGTGRRG
		İ				AGGGAHRRGSGRTGAREAGGR
						LRHGGLQVPSPARREGS*GPAS
		1				PAGCLAHL
14016	44384	Α	14098	346	957	ARYTLHIPTRLGSPAGFTQWIPH
						RGCRWSCLPVPRRALAFLSPW
						VVCSFPVDGTGRRGAGGGARR
						GGSGRTGAHGVGGR\PGMAGC
						RSRALPRGKAAKARREIERSAG
			}			G/TGTAAGPSTPSAAAGPGAKS
		ľ				PFARG/LAGPAAPSARPAKPTST
		ŀ				RNSSWPASAARSPGSRSWKMS
						LYMKKSASGQSQAQWIAKMW
						KSPRFKPLWLTS
14017	44385	Α	14099	124	663	SCLPVLRRALTFLSPWVVNGTG
						RRGVGGGARRGGSGRTGAHGV
						GGR\PGMAGCRSRALPRGKAA
						KARREIEHSAGG/TGTAGGPGT
						PSAATGPGAKSPIAPGQQGWLA
}						ALGINGAGERLFMKVAFSLKTC
						SLSCSYLRLTLSRGGSVHVLRG
						NLTVCKPQPALLRVVFHHQRQ
						ADCGALPVP
14018	44386	С	14100	1	672	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14019	44387	A	14101	3	1759	
14019	44307	^ <u>`</u>	14101]3	1739	SHGARRCLRKTAATSRCGGAP
		1				GP\RSRPPSTGQAAEAGDW/PAG CNP*GGLRAADGPGSRPPRGSI
						GLLCGQGGQGGSSPWVVHGTG
		1				RCGAGGGAPWGGSGRTGAHG
					:	
						VGGRLRHGGLQVPSPAPQEGS*
						GSARNRAQPRWAGTAGGPSTP
					:	SAATGPGRDGEGG\ETGAHVSA
		l				LKGTPGTEWPAHGLVPYSIGDQ
						TCLIFKVTFKAVVLKCGPQTTS
		l				STSISCPGSAN/SPPSPAG*/GSSG
						CGSSSLCSAHSVQEDSQPQ*KR
						TGLQS\TYGRSGSSSNEPRYHSD
						LPTSRSKLRNEDTFFGRRKKLS
[ļ	<u> </u>		<u>;</u>	EVQRTGQDPA*SHHTSSYTLPG
						L*TAEGKDRVENAA*KGSPNFQ
			ļ			RSGTFNPKGGARKTRQAWHPR
						APNCQAGTPPREGSSGPRTSLS
			1			AAFRIPQEVVNFLISGPITRAQR
	!					HGEPESPRGQTGSPPLGKTGSE
						GTDRKSTGADKKSTGDRQEVR
						EPTESPQGLDRKPTGMDRRSTG
		ļ				NRQKVRRQIGSPLGLNRKPNGT
						DKKSTRNRQEVRRRQSAGARQ
						ERAGSSTFLRLSALCGLLVDVN
						REQVAKQKWDLYSSRFTSGNR
14020	44388	Α	14102	1879	2607	GKPSLSLHPGHKCGISRCNGNV
		1			•	SSRLRSPAGFTQWIPHRGCRWS
		1				CLPAPRCALALLSPWVVDGTG
		1				RHGAGGGARRGGSGRTGAHG
		1		İ		VGGRLRHGGLQVPSPAPREGS*
						GPARNRAQRRWAGTAGGLSTP
						SAATGPGAKSPIARGQQGWPA
						APS\GGPPSPSPPGTPAGPQAPH
						AAPVPAR\PLPPHLPAS*GSGLQ
						PWPAQKGAPTVQGGG*RAPQM
						PPKWEPRQGRRIVAFTIEVGVL
14021	44389	Α	14103	2	272	GHWGMVNSPCRASATVSRGSP
						LPGVLPTSPPSATSTCRMRCHPR
1			1			RCPLPFRRLSPPSLTPP/PGKPPL
						LPRPHSMPGLGLPLGGLPVFHP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14022	44390	A	14104	101	1815	QSARQLRPCPGAPSLAGVENRR KKTTNLTSLAAPDTQSTCGQSC GCLHSMGSSCMQHSYYTGFLR SLWPCRCHRSAFSGQRRSEEKD SKSVRERSGRSQNVPARPALTS PLTQSPGPLAPGAGSWSRSQLG TGTGVLWWQRESATGDFRTES ARSSDPSRRDWPGAKRKREEG AATVLWACATASTAIICQDNAI SPLKSGNSPSNHLWRPQGQLPP VLGLAQPKRWRPNRLTTSPSPS RMMARPVDPQRSPDPTFRSSTR HSGKLEPMEATAHLLRKQCPSR LNSPAWEASGLHWSSLDSPVG\ PCRP*GLRPSTHGAR\TFSGARP GLGGYSPPPEEAMPFEFDQPAQ RGCSQLLLQVPDLAPGGPGAA GVPGAPPEEPQALRPAKAGSRG GYSPPPEETMPFELDGEGFGDD SPPPGLSRVIAQVDGSSQFAAV AASSHQCLSHHIPMSPSVGRNG LLTKPPSTSRRGKLTKRDDMYY FVPFRSLNDMMHMLPFIEDSR WTHNLDINREQKTNPLAGPKG RPISRTQAAASTSNSKTESQIQE
14023	44391	A	14105	39	877	KTEEAPFSLRYININNRSMDPLN AGPGPPRPLPPRSSEGPPGASISL CSEVPGPSCQGRFRGGRDAMS MPWREEERSLLAGE*PS*GG*S GRGVSRKCS*PGPRGVPRPRPL RSRPRAASSE\DQNSPLDQSLHR TRKTIPSSGGSTAIMPRRGSSSE TFFSAPSKTWVLGIPGETFISGP AAFP\APGGPRAAPRPSPSLNSL *SGGNGGPGPPRLHFSSCGKGP PRPSNSRN\GGQSACQGGSHLV C*SHRGSLGEGHAYQGDPCSPG HPAKAPSRAHAQARRSAHAAS LTSGPCDASCQAW
14024	44392	A	14106	358	546	CFPSRIRLALSRAPKWPAKRSPS EPEMS/PPVWSPHPDPNNTAPPK EQPPSSPGRVGRHGNWSV
14025	44393	Α	14107	22	549	DHCLCSVVRNKRDSVYPLPIPP CCKTSTISPP/LVDAVRP*DISVS KRVTSCAPLEPQPHPCLPTINAG R/TLARPQGAPTPNKEEGQAPG RATASGQAAPATTVLGDTTK/G PQGRHRGTSASAAGPAPFAVA DQLPPEDRAASAEGRK*AAAFP FCHRLVQDGGPPPPAAGVPGA GES

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
,				sequence		
14026	44394	Α	14108	146	2616	LAAVGMASHCFRQRRQSQSPA\
						GLRGAGSAYSPA\EGARSSHGQ/
						PPWRGRRRGPGDPVRGAARCR
						RPRAAAQPARPAARPPARPRPA
						P*AACCE/VPSAANSVAPSAAM
						*SSAGRDGASSQRSRRPHAAPP
						RSSALPANGK/PPSPVTRAARTC
						GARSSNTGA*SAARPKHSAASS
		l				RHPPPAA/MPLPARHA/PGGTSG
						RPLPTPQGSAPTSSASPRGPATP
						PPAGPRAPPPSPRRAAAPRPPGR
		1				QLSPAPGTPAAGGGGPPFPALR
						*SGSVFGRQLVGDGERGRACG
İ		1				ARGSAAVPALRTFVVSPSTVVA
						GAACPLAVALPAPDPLPCSE*A
		1				PLRSG*VPQPASLFCSQFRPQA
						WQHP*LLRAPCPRLGAAVPCV
		l				WCHSRALDSATPRWM*G*RLP
]				AVWL\PAFPSPPFNQAFVPPCQQ
		ĺ				NCSLQNRQTLSASAKIGTEEWD
						IAIKIPENVEVTLELGNEQRLEE
						FGGLRRRQEDEGKFGTPTDWL
		ł				NGCDQNADSNMDSEGQGEDG
						GCSFGGAGLRGSGWGDQTRET
		ŀ				FLALKANLRWPFRSSAQSQTSQ
1					İ	EGAAGARSGLKGGKKNPLKLP
		İ				KKQAKKMNKAFKQKQEKERK
		l				KHKELKAKASWKGLRPQVEFK
		1				NLAERRGEAAVPGCPVVAGHS
		1				WRLLSLARGCAKRARQLLFPSE
						KLLAALREGSTRGGGAASQPQ
		1				RKLRVCMRVLDSSVEDFHMDG
14027	44395	A	14109	1	885	MIISIDTEKALDKILHPFILKTLN
						KLGIDGTYLKIIRALYDKPTANI
						IMNEQKLEAFPLKSNTKQGCPF
						LPLLFNIVLEVLARAIKQEKEIK
						SIQIGREEVKLSLFADDMIVYLE
						NPIISAPNLLKLISNFSKVSGYKI
						NVQKSQAFLYTSNRQTEIQMM
]						SELPFTIATKGIKYLGI\HPLTRD
						VKDLFKENSKPLL/KELEKTTLN
						FIWNQKRAHIAKTMLSKKNKA
						GGVMLPDFKLYYKAAVTKTA
						WLGDDQVFLIPRGHISDYHMG
						KNLGQYPAFQDRGPCGFPQCIV
						PLVYRD
	1	Ь	l	<u> </u>	L	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide inscrtion)
				sequence		
14028	44396	A	14110	453	891	LIHLSTDALRVSGWGTVRSFSS
						HEAIFQTIAWGETLDNTLLSRA
						EVPAAFHGALCPWFIETREWQ*
						LLQSILLVNIWLTRHVLHSPRSL
					!	KP*FYTT*VFVCSKLGQSGWGK
						VAGAKLQMNNISAKQLFKVQV
						LFKIESLMSSLST
14029	44397	A	14111	452	2922	KMGLAREWILRFLVLILQEIRS
						MRQKENKDIQDLNSALHQADL
						IDIYRTLHPKSTEYTFFSAPHCI/
						RTYSKIDHTLGSKALLSKCKITE
						IITD\$QTTVLSELRIKKLTQ\NRS
						ATWKLNS/YLNDYWVHNEMK
						AEIKMFFETNENKDTTYQNLW
						DTFKAVCRGKFIALNAHKRKQ
						ERSKIDTLMSQLKELEKQEQTH
	ł					SKVSRRPITG\SEIEAIINSFTNPK
						RVPGPDGFDEPNFYQRYKQEL
						VPFLLKLFQSIEKDGILPN*FYE
						ASIILIPKPGRGTTKK/EENFRPIS
						LMNIDAKILNKILANRIQQHIKK
						LLHHDQVGFIPGMQGWFNICKS
						INVIQHINRTSDKNHTIISIDAEK
						AFNKIQQLFMLKTLNKLGTNG
						MYLKIVRAIYDKPTANIILNGQ
					·	KLEAFPLKTGTRQGCPLSPLLF
						NIVLEVLARAIRQEKEIQGIQLG
						KEEVKLSLFADDMIVYLENPIV
						SAQNLLKLISNFSKVSGYKINLQ
						KSQAFLYTNNRQTESQIMSELP
						FTTASKRIKYLGIQLTRDVKDLF
	i					KENYKPLLNEIKEDTNKWRNIP
						CSWVGRINIVKMAILPKVTYRF
						NAIPIKLPMTFFTELEKTILKFR
İ						WNQKRAHIAKTILSQKNKAGGI
						RLPDFKLFYKATVTKTARYWY
						ENRDIDQWNRTEPLEIMPHIYN
İ						HLIFDKPDKNKQWGKDSLFNK
						WCWENWLAICRKLKLDPFLTS
14030	44398	Α	14112	3	3349	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14031	44399	A	14113	294	1107	RRHRLENYIPRTAQSPHPRKAL
				-		ARSESKRDGGFKNNWSFDHEE
						ESEGDTDKEPGPLDAPPRP/ESG
						PRAPGKGL*RRSRSPGPSERGSA
						ARLTAEPEARNPAASTLLRRGP
		İ				SSRRARRRCGPWIKKEAAWEE
						EDGGRQERRLGQDLRTRTLNK
			ļ			SKGGIDSRIISLGQHSHPTHAKL
						WLDQSLREMEVLKIIGALIMKK
						KVKEIQIKSRARSTPLHDQIGTT
						SPRERTIASKRHPYYITPKGIRY
						EQDTLSRTPITTIRSRQRPLGPLT
						ASPPPSFPTLAPPTQAIKPSLHLR
				1		EGKPTPTPGERSTRPHHPEPPRK
						EYALSSLYPPTYLLAPTSTDLG
						NAHPTQEPQFNVDATHPRQNR
						FIPVAGNSPKRSSQFPIVFAPKK
						NTTPPYLDFAAILGLHRRPPRSR
						FRTSYPPKEFRSSPPERTAPRN
14032	44400	Α	14114	2	297	VQKSVLCSFQAGTMEGVEEKK
						VPAVPETLKKKRRNFADAIQFL
						QTVFLAF*YTLLKGRQ\VYWLV
					:	EE*HQFYKISYRKKMVQRIRQH
						PQSCTVQKSWWP
14033	44401	Ã	14115	3	754	GTMEGVEEKKKEVPAVPETLK
						KKRRNFAELKIKRLRKKFAQK
						MLRKARRKLIYEKAKHYHKEY
						RQMYRTEIRMARMARKAGNF
						YVPAEPKLAFVIRIRGINGVSPK
						VRKVLQLLRLRQIFN\GTF\VKL
						NKASINMLRIVEPYIA*EYPNLK
						SVNELIYKHGYGKINKK*VALT
						DNTLIARPLGKYGITCMEDL/IH
						EIYTVGKCFKEANNFLWPFKLS
						SALSGMK/KKTTHFVEGGDGG
						NRVDQINTLNRRMN
14034	44402	Α	14116	1	388	

NOC.	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
14035	_	•				5	*=Stop codon, /=possible nucleotide
14035		sequence		09/540,217	1	of peptide sequence	deletion, \=possible nucleotide insertion)
MEPOPYNSKELNSANNQCLLE KPKLRQRLKDKEYDQGHTAVQ CQSWYYVLTPDSFCQQLLLPS TENFHECSRSDTAGHLRTGHVA SNWLSEHSPOTMPAAMQMOK KMQNTSSWFPQKAPEKVKPMS KLWQTPSQNDTLYKAVKALTL SSPERNSITINKKDVHTETSKEY SSPERNSITINKKDVHTETSKEY KAERSQNQNASSPRNINSSE ARERNWSENSPEDLTEVVFRR WYITNSSELKEHVLTQCKEAKN LEKRLEELLTRITGLEINQAERI LEKRLEELLTRITGLEINQAERI SETEDQLNEIKHEDKIREKRMK RNEQILQEIWAYVKIREKRLIG VPENDREINGKLENTLQDTIQE NFPELAR*ANIQIQ NFPELAR*AN					sequence		
MEPOPYNSKELNSANNQCLLE KPKLRQRLKDKEYDQGHTAVQ CQSWYYVLTPDSFCQQLLLPS TENFHECSRSDTAGHLRTGHVA SNWLSEHSPOTMPAAMQMOK KMQNTSSWFPQKAPEKVKPMS KLWQTPSQNDTLYKAVKALTL SSPERNSITINKKDVHTETSKEY SSPERNSITINKKDVHTETSKEY KAERSQNQNASSPRNINSSE ARERNWSENSPEDLTEVVFRR WYITNSSELKEHVLTQCKEAKN LEKRLEELLTRITGLEINQAERI LEKRLEELLTRITGLEINQAERI SETEDQLNEIKHEDKIREKRMK RNEQILQEIWAYVKIREKRLIG VPENDREINGKLENTLQDTIQE NFPELAR*ANIQIQ NFPELAR*AN	14035	44403	I I	14117	<u> </u>	11145	MTRNYEWI I FAFSSDODIA FOU
	14033	44403	^	1411/	1	1143	
CQSWYYLFPDSFCQQLLLPS TENFHECRSDTAGHLRTGHVA SNWLSEHSPQTMPAAMQWDK KMQNTSSWFPQKAPEKVRMS KLWQTPSQNDTLYKAVKALTL SSPERNSINTNKKDVHTETSSK GHQHQRFKVDKSMKMRKNQH KKAENSQNQMASSPPRNINSSPARERNWSENEFDELTEVVFRR WVITNSSELKEHVLTQCKEAKN LERRLEELLTRITIGLEINQAEEN SETEDQLNEIKHEDKIREKRMK RNEQILQEIWAYVK:RPKLRLIG VPENDRENONKLENTLQDTIQE NFPELAR*ANIQIQ NFPEL							
TENFHECSRSDTAGHLATGHVA							
SNWLSEHSPQTMPAAMQWDK KMQNTSSWFPQKAPEKVRPMS KLWQTPSQNTLYKAVALTL SSPERNSINTNKKDVHTETSSK GHQHQRPKVDKSMKMRKNQH KKAENSQNQNASSPPRNINSSP ARERNWSENEFDEL TEVVFRR WVITNSSELKEHVLTQCKEAKN LEKRLEELLTRITGLEINQAEERI SETEDQLNEIKHEDKIREKRMK RNEQILQEIWAYVK:RPKLRLIG VPENDRENORKLENTLQDTIQE NFPELAR*ANIQIQ			1				
KMQNTSSWFPQKAPEKVKPMS KLWQTPSQNDITYKAVKALTI SSPERNSINTNKKDVHTETSSK GHQHQRPKVDKSMKMRKNQH KKAENSQNQNASSPRNHNSSP ARERNWSENFEDELTEVVFRR WVITNSSELKEHVLTQCKEAKN LEKRLEELLTRITGLEINQAEER SETEDQL.NEIKHEDKIREKRMK RNEQIL.QEIWAYVKRPKLR.LIG VPENDRENGKLENTLQDTIQE NFPELAR*ANIQIQ NFPE							1
KLWQTPSQNDTLYKAVKALTL SSPERNSINTIKKDVHTETSSK GHQHQRPKVDKSKMKRKNQH KKAENSQNQNASSPPRNHNSSP ARERNWSENEPDELTEVYERR WVITNSSELKEHVLTQCKEAKN LEKRLEELLTRITGLEINQAEERI SETEDQLNEIKHEDKIREKRMK RNEQIL QEIWAYVKREVLRLIG VPENDRENGKLENTLQDTIQE NFPELAR*ANIQIQ NFPELAR*ANIQIQ NFPELAR*ANIQIQ NFPELAR*ANIQIQ WPENDRENGKLENTLQDTIQE NFPELAR*ANIQIQ WPENDRENGKLENTLQDTIQE NFPELAR*ANIQIQ WPENDRENGKLENTLQDTIQE NFPELAR*ANIQIQ NFPELAR*ANIQIQ WPENDRENGKLENTLQDTIQE NFPELAR*ANIQIQ WPENDRENGKLENTLQDTIQE SETEDQLNEIKHEDKLENTLQDTIQE NFPELAR*ANIQIQ WPENDRENGKENTLQDTIQE WPEND	l						-
SSPERNSINTNKKDWHTETSSK GHQHQRPKVDKSMKMRKNQ KKAENSQNQNASSPRNHNSSP ARERNWSENEFDELTEVVFRR WVITNSSELKEHVLTQCKEAM LEKRLEELLTRITGLEINQAEER SETEDQLNEIKHEDKIREKRMK RNEQILQEIWAYVKRPKLRLIG VPENDRENGNKLENTLQDTIQE NFPELAR*ANIQIQ NFPELA				•			
GHQHQRPKVDKSMKMRKNQH KKAENSQNQNASSPPRNHNSSP ARBRIWSENEPGELTEVVPRR ARBRIWSENEPGELTEVVPRR WVITNSSELKEHVLTQCKEAKN LEKRLEELLTRITGLEINQAEERI SETEDQLNEIKHEPDKIREKRIMK RNEQILQEIWAYVKVRPKLRLIG VPENDRENGRKLENTLQDTIQE NFPELAR*ANIQIQ							1
KKAENSQNQNASSPPRNHNSSP							
ARERNWSENEFDELTEVVFRR WYITNSSELKEHVLTQCKEAKN LEKRLEELLTRITGLEINQAEERI SETEDQLNEIKHEDKIREKRMK RNEQILQEIWAYVK\RPKLRI.IG VPENDRENGKLENTLQDTIQE NFPELAR*ANIQIQ NFPEL							1 7 7
WVITNSSELKEHVLTQCKEAKN	ŀ		İ				1
LEKRLEELLTRITGLEINQAEERI SETEDQL,NEIKHEDKIREKRMK RNEQILQEIWAYVK\RPKLRIG VPENDRENGNKLENTLQDTIQE NFPELAR*ANIQIQ 14036							i i
SETEDQLNEIKHEDKIRERRMK							1
RNEQILQEIWAYVK\RPKLRLIG			1				1
Table			İ				1
14036 44404 C	1			,			1 1
14036							1 ' ' 1
14037	14036	44404	C	14118	291	551	INTELAK ANIQIQ
14038				<u> </u>			
PPQPSQ/CKQWGDPTPVEAC/CV SCPTQGFISPTRGGPPCKHESLV VPLVGTPRRNQPGGHIQHSQR WAPVLT					1.		ITI SSPPHSEIGHKTPASGPPOKI
SCPTQGFISPTRGGPPCKHESLV	114030	1		14120	102	307	1
VPLVGTPRRNQPGGHIQHSQR WAPVLT							1 1 1
WAPVLT			ĺ				1
14039			ŀ				1
RSMRC/GP*GF\$PGAEA*TEP/C	14039	44407	A	14121	999	2661	
HYGGRGFPCGPFSLGCLAT\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	111037	1	` `	1 1123		2001	- I
WLPGPRPPCSPWAGP\CGLCPLP AAAAERQPALSVPGASPSCRLL PLVVLCHGSHLNGGRVLLRPG GSPCGHCHNCAGARRGLGHVL QGGAQARQREQLQR 14040			ł				
AAAAERQPALSVPGASPSCRLL PLVVLCHGSHLNGGRVLLRPG GSPCGHCHNCAGARRGLGHVL QGGAQARQREQLQR 14040							
PLVVLCHGSHLNGGRVLLRPG GSPCGHCHNCAGARRGLGHVL QGGAQARQREQLQR 14040			1				i
GSPCGHCHNCAGARRGLGHVL QGGAQARQREQLQR 14040							1
QGGAQARQREQLQR							i i
14040							l i
14041	14040	44408	В	14122	164	1042	
14042		1	В		I	566	
TVPSQLVQYCPSF*ISWKPMVY QKVPA*PRPPRTPDHHPDPGPA PHPRQHHPPPRSPDAEP/SPTTPT RRRPPTQPPDPHPKRPPPHSKPH PPQQPPTPTATPTEQQRHTAPP TPRPPTTHPPSTPKHPA\PHTAPP PTKNTKPPTPQRTTQDKGRQKK PRRQPHHSEPNKKSPKAHI 14044 44412 C 14126 I 1128 14045 44413 A 14127 319 536 14046 44414 C 14128 231 425 14047 44415 B 14129 I 1431 14048 44416 A 14130 75 393	14042	44410	В		1539	1674	
QKVPA*PRPPRTPDHHPDPGPA		1	A		 	780	QKAKKGKLKRSKQNSFPGLKK
QKVPA*PRPPRTPDHHPDPGPA							,
PHPRQHHPPRSPDAEP/SPTTPT RRRPPTQPPDPHPKRPPPHSKPH PPQQPPTPTATPTEQQRHTAPP TPRPPTTHPPSTPKHPA\PHTAPP PTKNTKPPTPQRTTQDKGRQKK PRRQPHHSEPNKKSPKAHI 14044 44412 C 14126 I 1128 14045 44413 A 14127 319 536 14046 44414 C 14128 231 425 14047 44415 B 14129 I 1431 14048 44416 A 14130 75 393 393			Ī				
RRRPPTQPPDPHPKRPPPHSKPH PPQQPPTPTATPTEQQRHTAPP TPRPPTTHPPSTPKHPA\PHTAPP PTKNTKPPTPQRTTQDKGRQKK PRRQPHHSEPNKKSPKAHI 14044 44412 C 14126 I 1128 14045 44413 A 14127 319 536 14046 44414 C 14128 231 425 14047 44415 B 14129 I 1431 14048 44416 A 14130 75 393 393							-
PPQQPTTPTATPTEQQRHTAPP TPRPPTTHPPSTPKHPA\PHTAPP TPRPPTTHPPSTPKHPA\PHTAPP PTKNTKPPTPQRTTQDKGRQKK PRRQPHHSEPNKKSPKAHI 14044 44412 C 14126 I 1128 14045 44413 A 14127 319 536 14046 44414 C 14128 231 425 14047 44415 B 14129 I 1431 14048 44416 A 14130 75 393 393 14048 14130 75 393 14150 1415							
TPRPPTTHPPSTPKHPA\PHTAPP PTKNTKPPTPQRTTQDKGRQKK PRRQPHHSEPNKKSPKAHI 14044 44412 C 14126 I 1128 14045 44413 A 14127 319 536 14046 44414 C 14128 231 425 14047 44415 B 14129 I 1431 14048 44416 A 14130 75 393	}		I				-
PTKNTKPPTPQRTTQDKGRQKK PRRQPHHSEPNKKSPKAHI							
PRRQPHHSEPNKKSPKAHI							PTKNTKPPTPORTTODKGROKK
14044 44412 C 14126 I 1128 14045 44413 A 14127 319 536 14046 44414 C 14128 231 425 14047 44415 B 14129 I 1431 14048 44416 A 14130 75 393	}						
14046 44414 C 14128 231 425 14047 44415 B 14129 1 1431 14048 44416 A 14130 75 393	14044	44412	C	14126	1	1128	
14047 44415 B 14129 1 1431 14048 44416 A 14130 75 393	14045	44413	Α	14127	319	536	
14048 44416 A 14130 75 393	14046	44414	С	14128	231	425	
	14047	44415	В	14129	1	1431	
14049 44417 C 14131 174 371	14048	44416	Α	14130	75	393	
	14049	44417	С	14131	174	371	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14050	44418	Α	14132	79	164	
14051	44419	Α	14133	343	775	
14052	44420	Α	14134	554	724	SEGPSKEPTHRRMKFSHHDDLI
						FLDLSN*QPPNSPEPHPPQSP*KP WPQNPSSRQM
14053	44421	Α	14135	20	242	VSFLSMGSGHCIRSTRGSKMVS
]	WSVIAKIQEI*CEEDERKMARES
					}	\LAESMSTYVMMNHTYDS*GK
						THILIMKK
14054	44422	Α	14136	229	366	
14055	44423	Α	14137	131	505	
14056	44424	Α	14138	1	1317	AGCSCGSRAMAEEQGRERDSV
						PKPSVLFLHPDLGVGGAERLVL
						DAALALQARGCSVKIWTAHYD
		}				PGHCFAESRELPVRCAGDWLPR
		ŀ				GLGWGGRGAAV\CAYVRMVFL
		i				ALYVLFLADEEFDVVVCDQVS
ļ						ACIPVFRLARRRKKILF\YCHFA
		ŀ				DLLLTKRDSFLKRLYRAPIDWI
						EEYTTGMADCILVNSQFTAAVF
		ŀ				KETFKSLSH\KDPDVLYP\SLNV
İ						TSFDFSCS*KAGMT*SPRGKNS
		ŀ				WLLSINRY\EREGKIWTLGTGK
		ŀ				ALVQLRGRLTSQDWERVHLIV
						AGGYDERVLENVEHYQELKK
						MVQQSDLGQYVTFLRSFSDKQ
ļ						KISLL/RQLARVCFYTPRQ*GTL
						GIVPLGRPCYMAVPQFICCLIRV
						GPFGSSI*PQCSQGFCVEPDPVH
						FSEAIEKFIREPSLKATMGLGWE
						EPE*REKFSPGSILQEQLYRYVY
14057	44425	Α	14139	1	648	MESKEEQGGVADHQKATRGIP
						TPQPREVVSERATQRGNRAFST
	į					KLCNPKTNRKQQQQYQQQK
			1			GRTKTLPKGQHPQRSKLDKLTE
1						MRKHQQKNTKNPKGQSPCSPP
-						NDCNVSPARVHNWTGDEMEEL
						TEVGFKKWVRKNCAELKEHVL
		1				TQCKEAKNLDKRLEELLTRKTS
						LERNINDLMELKT/RA*ELCKA
-						YTSINS*IDQAEERISEFEDHLAE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14058	44426	Α	14140	1	1365	MPVKGGTKCIKYLLFGFNFIFW
						VSERDCRALLSGPTCSRAPDTG
		1				RGRECRQLALPAPGRHRAGRES
						ALRLPAGSKAGSRAGRDGRDG
						RIRVGVSGTLFGSTTGLFNQSL
						AETGFTPYTQKSNDFLVKSLKI
						WHTYFKNFHQEARLYIRTSPGS
						NWNWFCTLGIESPEQPGVYEG
						QRPRVVTKSFASLTLKRVGTVL
						LLAARINADVLYVLQLAGIAVL
						AIGLWLRFDSQTKSIFEQETNN
						NNSSFYTGVYILIGAGALMMLV
						GFLGCCGAVQESQCMLGLFFG
						FLLVIFAIEIAAAIWGYSHKDEV
						IKEVQEFYKDTYNKLKTKDEPQ
						RETLKAIHYAVCRLGKDTLLRF
l						LRIVSAHRLNCCGLAGGVEQFI
						SDICPKKDVLETFTVKSCPDAIK
		ŀ				EVFDNKFHIIGAVGIGIAVVMPS
						FRSNHPDPHVSSYLL\IFGMIFS
		l				MILCCAIRRNREMV
14059	44427	В	14141	98	1064	
14060	44428	Α	14142	560	964	KRLPRILGIRGKPCCPDTSRAGR
						RVRGRAAAPCREAARGRGQRR
	ŀ		1			FLPPTWRCETGAATMFPSPALT
						PTPFSVKDILNITVRPRS/LGQAA
						LTKRLSASL\SPQSCARCRRRWS
						CVILSAVRGN*YAGSRAHRDCT
						Q
14061	44429	С	14143	189	452	
14062	44430	Α	14144	1	647	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	i e	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14063	44431	A	14145	890	2905	LRLSLKFPMSQWTPEYNELYTL
						KVDMKSEIPSDAPKTQESLKGI
						LLHPEPIGAAKSFPAGVEMINSK
						VGNEFSHLCDDSQKQEKEMNG
						NQQEQEKSLVVRKKRKSQQAG
						PSYVQNCVKENQGILGLRQHL
						GTPSDEDNDSSFSDCLSSPSSSL
						HFGDSDTVTSDEDKEVSVRHSQ
		:			<u> </u>	TILNAKSRSHSARSHKWPRTET
						ESVSGLLMKRPCLHGSSLRRLP
						CRKRFVKNNSSQRTQKQKERIL
						MQRKKREVLARRKYALLPSSSS
					.	SSENDLSSESSSSSSTEGEEDLF
						VSASENHQNNPAVPSVTYGQP
						WSKNIKEKTLEINSSQALTAYE
						CLHSAHVINLISPFYHHFPSSQE
						RSYLSTLSTLSNKKTGSVWFTR
						SPKVTQFFPGGAITGTLSQRPER
						EILRPLALPGKGNTPPYFCSPSM
						GCTHCPTSPNEMNQGISEDTAA
						APVRDIYIKPPSPWDRAPGGRG
						GCGHSFSRLKSPYLRALKSAAD
						FPAQHSSSDKGQAASSSGSLTP
						VYPVWVTPPSRGRQTPHRGEL
						WLASDGCPSGRKLPEEGTGSNL
						CCSAASAGDTQANRDHNSSPA
						REQTWTENKFDKLTEAGLRRW
						VINSSELKEPVLTQCKEAKNLE
						KRLDESPTRITSSEKNINDLMEL
						KNTAQELHEEHTSINSQ/DQAE*
14064	44432	Α	14146	1	737	LNRGEQRAVRYYSHMKLNMA
						EEEDYMSGFLH*CPRRYQTRIA
						NAKANPRSPSKR\EKQQEANLK
						NRQKSLKEEEQERRDIGLKNAL
						GCENKGFALLQKMGYKSGQAL
						GKSGKSGIGHEASLKRKAEEKL
						ESYRKKIHMKNQAEEKAAEQF
						RMRLKNKQDEMKLEGDLRRSQ
						RACQQLDVQKNIQVPREAWY
						WLRLEEETEEDEEEKEQDEDEY
						KSEDLSIKKTYLQIVQDQLLQI
						MTKIIPNKVET
14065	44433	A	14147	1	1860	
14066	44434	A	14148	1	1203	
14067	44435	В	14149	1	1119	
14068	44436	A	14150	1	1407	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14069	44437	A	14151		1731	ERSSPATEQSWMENDFDELRE EGFRRSNFSELKEKVRTHHKQD KNLEKRSWFFEKINKIDRPLAR LIKKKREKSQINAIKNDKGDITA DPTEIQTTITEYYKHLYANKLE NLEAMDKFLDTYTLPRINQEEV ESLNR/LNNRL*N*GNN**LTNQ K/NVQDQMD*QLNSTRGTRRS WY/RFLLKLFQSIEKERILPNSFY EGIILLIPKPGRDNTKKDNFKPIS LMNMDAKILNKILANRIQQHIK KLIHHDQVGFIPGMQGWFNICK SINVIQHINRTKDKNHMIISTDA GKAFDKIQQPFMLKTLNKLGID GMNLKIIRAIYDKPTANIILNRQ KLEAFPLKTGTRQGCPLSPLLF NIVLEVLARAIRQEKEIKGIQLG KEEVKLSLFADDMIVYLENPIV SAQNLLKLISNFSKVSGYKISVQ KSQAFLYTNNRQTESQIMSEIPF PIASKRIKYLGIQLTRDVKDLFK ENCKPLLKEIKENTNKWKNIPC SWIERINIVKMAILPKVIYSFNAI PIKHPMTFFTELEKTTLKFIWNQ KRARITKSILRQKNKAGGITLPE
	<u> </u>					FKLYYKTTVTKIAWYG
14070	44438	В	14152	1	906	
14071	44439	A	14153	3	553	EHSSSPATEQSWMENDFDELRE EGFRRSNYSELKEEVRTHGKEV KNLEKKLDEWLTRITNAEKSLK DLMELKTTAREICDECTSLSSR CDQLPRPDGFTAEFYERYKEEM VPFLLKLFQSIEKE/VNPP*LTL* GQHHPDTKAWQRHNKKREF*T NIPDEH*CKNPQ*NTGKPNPAA HQKAYPP
14073	44441	A	14155		969	MKAEIKMFFETNENKDTTYQN LWDTFKAVCRGKFIAVNVHKR KQERSKIDTLTSQLKELEKQEQ THSKASRRQEITKIRAELKEIET QNTLQKINASRSWFFERTNKID RPLARLIKKKREKNQIDAIKND KGDITTDPTEIQTTIREYYKHLY ANKLENLEEMDKFLDTYTLPRL NQEEVESLNRPITGAEIEAIINSL PTKKSPGPDGFTAEFYQRYKEE LECSSSPAMEQSWTENDFDDLR EEGFRRSNFSELKEEVQTHRKE AKNLEKRLEKWLT\RITNVEKS LNYLRELKTMARELC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14074	44442	A	14156	377	687	LLEGKLTNRKDIHTKTPSV/SPP
						SSKTKEHSSSPAMEQSWMEND
						FHELREEGFRRSNFSKLKEEVR
						THRKEVENLEKRLDEWLTRITK
						VEKALNDLMELKTMARE
14075	44443	Α	14157	1	711	
14076	44444	Α	14158	1	831	
14077	44445	A	14159	1	744	
14078	44446	В	14160	1	1479	
14079	44447	Α	14161	1	810	
14080	44448	Α	14162	3	651	
14081	44449	Α	14163	2	1067	
14082	44450	Α	14164	2	2063	
14083	44451	Α	14165	1328	1639	PTNAHETSGRYIAGSSEKRTSA
1						RTCGGTSAAARPAGTSMQTRC
						RQPPPSS*HAVPRVAQSSDSGL
<u> </u>						KATHPQAYQPCGHTPTRLPQES
<u> </u>		i				NCIEHERGLGKKKKKKK
14084	44452	Α	14166	1	620	MTLGYGQRGANKDLRTGFDPD
						LCTMDNFAEGDFTVADYALLE
		1				DCPHVDDCVFAAEFMSNDYVR
		1				VTQLYCDGVNDSFLIGGLLRIG
		1				CKIENERSSSPAMEQSWTENDF
		İ				DELREEGFRRSNYSELKEEVR\T
		ŀ				NGKEVRNFEKKLDEWITRITNA
		ŀ				EKSLKDLMELKTTARELCDKC
	1	İ				TNLSNRCDQLEERVSAMEDEM
						NEMKHEDKFREKE
14085	44453	Α	14167	1	514	MGKKQNRKTENSKNQRTSSSP
		l				KERSSSPAAEQSWMENDFDEL
		İ				REEGFRRSNFSELKEEVRTHGK
		l		ŀ		EVKNLEKRLDKWLTRITNTQKS
}				•		LKDLMELKTTARELHDECTSLT
ļ						NQFDQLEERINNI\EDH*QD**R
						RKERRIK*MQ*KMIKGISPLIPQ
						KYKLPSENTINTSMQIN
14086	44454	Α	14168	1	366	MDTFLQAERKDYMEAYELIEQ
}						EEQGEREPVAVNNILSTEALMA
						NAKSTLMFYAVGFLWQPGGGR
						GHKAASNPGVTRDVVGSLSSLT
					:	FLRGERSSSPAMEQSWTDNDFD
						EL\REEGFR*SNFS
14087	44455	A	14169	1	1026	
14088	44456	Α	14170	1	3906	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14089	44457	A	14171	1643	2721	I IKPTKMGKKQSRKTGNSEKQST
14009	44437	^	14,7,1	1043	2721	SPPPKERSSSPAMEQSWMENDF
		l				DELTGAGFRRLVITDFSELKED
						VQTHHKEAKNLEKRLDKCRTN
						1 -
						G*LEWMNG/ITRITNAEKSLKDL
ŀ				ľ		MELKTKARELHDECTSLRSRFD
						QLEERVSVMEDQKK\MNEMKR
	·	1				EGKFREKRVKRNEQSLQEIWD
		1	ľ		l.	YVKRPNLRLIGVPESDGENGTK
		İ				LENTLQDIIQENFPNLARQANV
	1	ŀ	·			QIQEIQRTPQRYSSRRATPRHIIV
		1				RFTKVEMKEKMLRAAREKGQ
	1					VTLKGKHIRLTVDLSAETLQAR
	1					REWGPIFNILKEKNFQPRISYPG
						KLSFISEGEIKYFTDKQMLRDFV
		1				TTRPALKELLKEVLNMERNNR
		<u> </u>				YQPLQNHAKM
14090	44458	Α	14172	351	898	SGVTAPWRPNGTLLITAATASH
		1				PVLAIPAHLCSPRCPSFLTTVTC
						SGPQHHAARGRVPFHC/SECGK
						SFRYRSDLRRHF/ARHTALKPH
						ACPRCG/KGFKHSFNLANHLRS
ļ						HTG/ERPYRCSACPKGFRDST/G
						LLHHQVKTITT*PD/CGGCLVSN
						IVGSYSRPPWPCTRGPRRPLGSP
						SPAPGSVCLC
14091	44459	Α	14173	1	551	
14092	44460	Α	14174	3031	3207	TIHGCWWDPLSSIF*PFWGRGFI
						SHKICQAQISFRVISGTVLSLDK
		ı	}			LLIMFHKAIFASP
14093	44461	Α	14175	3555	3957	ENGQKTWTGTSQRRYTGDRM
						AQIQNPDTKCW*ECKATGTRIH
						CWWECKMVQPLWKS\WQFLT
		1				KLSILLPYNLAIILLGNYPNALK
:						MYAHTKGFTCMFIAASFIIAKT
}						WKQPRRPSVGEW/INNL*LVQT
		İ				MEYYSAL
14094	44462	A	14176	3	883	CQPYPRPRPRTGTHGSCQPPSQ
						GSAEP*G*CLPTLLRMALKAQP
						TKVNPRTLLDSDPDVC/S*AAHS
						RVLITSPSRAPWRLPSP*RRCPA
			į			GTCMPSTATSAR*CSQRPGSSW
						*T*TGSSLGSQACGPSSSTAPPSS
						SWSACTCGCAAAARCSCAASS
						TRSGPTCGSSPPASSCASSTPAP
						GTTPSSTLTSWASSPWSTPCPGS
ĺ						AGPSSWSSSSSATPSASASTRTL
						SPGSPAAP*PWPTALSRLTEQEA
						VWGA WGISHGNSWDKETKLV
						RLPHPCSKSPCFRPSLQPYRAHR
L	<u> </u>	<u> </u>	<u> </u>	<u> </u>	L	AP

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14095	44463	A	14177	3	395	SAEVGAAETTLTELRRTVQSLEI DLDSMRNLKASLENSL\GILLHL ESELAQTRAEGQRQAQEYEAL LNIKVKLEAEIATYRRLLEDGE DFNLGDALDSSNSMQTIQKTTT
14096	44464	A	14178	2	1044	RRIVDGKVVSETNDTKVLRH AKTSSQMPSPQEGMCGKACTP ALSQADSLCPLLRLASEVEGYS
						LPACAEPYVQSECLSHLSVWSL QHALSSLASMSFTTCSAFTNYW SPGSVQVPSYGTQPVSHAASVY
						AGLGGSGSRISVSHFTNG\GGLA GMGGIQNEKETMQSLRDRLAS YLDRVRGLETENWKLESKIQEH
						LNVTRL*LETEIEALKKELLFM KKNNEEEAKGLQAQIASSGLTV EVDAPKS/QDLAKNMADSRAQ
						YDKLARKNREELDKYWSQQIE ESTRVVTTQSAKVGAAEMTLT
						ELRHRVQSLEIDLDSTRNLKAS LENSLREVEARYALQMEPSLLF SKIIFGRVWEISYLHVFEGKQV

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14097	44465	Α	14179	1	2040	MPVPQLPPPVSRLAATAAASVT
		l				EPGPNPEAVRSASAEPEVKEGF
		l				PSQNHSVGHSNLIPSGGSVAPSS
		l				GRSHYGYASEDGTCSKLRVRPR
		l				LAAAPAAPFAFVADLASCCQG
		1	}			LSFFEGILLVRGPYRRLRATVV
		l				RDLVAVRMAEEQEFTQLCKLP
						AQPSHPHCVNNTYRSAQHSQA
						LLRGLLALRDSGILFDVVLVVE
		1				GRHIEAHRILLAASCDYFSMSFT
						TRSTFSTNYWSLGSVLATTYGA
		1				RLVSSTARVYAGAGGSGSRISV
						SRSTSFQGGLESGGLAAGMAG
		1				GLAGMGGIQNEKETMQSLNDR
						LASYLDRVRSLETENRRLESKIR
						EHLEKKGPQVRDWSHYFKIIED
						LRAQIFANTVDNARIVLQIDNA
		ĺ				RLAADDFRVKYETELAMRQSV
						ENDIHGLRKVIDDTNITRLQLET
						EIEALK*ELLFMKKTPEEEVKG
						LQAQIASSG\LTVEV\DAPKSQD
		l				LAKIMADIR\AQYDELARK\NRE
		ļ				ELDKYWSQQIEESTTVVTTQSA
		1				EVGAAET\TLTELRRTVQVFGD
		ł				STWTSMRNLKA\NLENSL\GEV
		1				EARYAL\QMEQL\NGILLHLESQ
						LGQTPRTEAQRQAQE\YEALL\N
						IKVKL\EAEICHLTRPPSWKIGE
						DFNL\GDS\LDERNSMQTIQKTT
		1				TRRISWIGQSGVLRPIDTKVLEA
						LSQQEASGTLLGKQEANKKFQS
14098	44466	A	14180	275	550	
14099	44467	В	14181	729	909	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14100	44468	Α	14182	1	1415	MGTQCDLLAKSLEDCPVDEVE
						DAFQGLGKEDEEIEQFSDKTFG
						SGAIDDDWREAHECLAELEVK
						QPVAVIEQTGNGERDEMDLLD
		·				DHEENLAERLSKMVIENELEDP
						AIMRAVQTRPVLQPQPGSLNSSI
						WDGS/VSSEANPRTTACSGNAY
						SVCIRICFASEA/SPRVQKMIGTF
		•				LNEHYQGGQLHLSLAVLLLELS
		·				P*APHLNRWLYPASPNRFCVRS
						LSMFGP/QMPPRYPAPYGERMS
						PNQLCSVPNSSLLGHPFPPSVPP
						VLSPLQRAQLLGGAQLQPGRM
						SPSQFARVPGFVGSPLAAMNPK
						SQAPMFRPDTTHLHPQHRRLLH
						QRQQQNRSQHRNLNGAGDRGS
						HRSSHQDHLRKDP\YANLMLQ
		1				REKDWVSKIQMMQLQSTDPYL
						DDFYYQNYFEKLEKLSAAEEIQ
						GDGPKKERTKLITPQVAKLEHA
						YKPETIICQALCRALHINYLTKA
						FQQAFEVSASIITIYR
14101	44469	A	14183	204	560	QGYWRGCTSSGGDGGCRGCW
						SPGKRR*GLSRTASGPPAAVAS
	1	l				PGRPRSPPSARTG*S*PGA*ARP
						CSRTW/PPPRCPLRPPSAARASA
						SRV/APL/GEPAPANTNIYCAAL
						THRATKSSTREF

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14102	44470	Α	14184	59	1695	DMPWWPVLPWIKDSSPSSLEM
						NPDNTSGLRQKSVEAEPCSLW
					İ	MVHRYTPQSTGEACTMAKAA
						GPSSP\GSCPPRPLGADPWPMSR
	1					LSH*EC*KTGTTASGYRRHQET
						APPESSGSPVGLSAGLWLSWA
						WPVSPACLHPGPPLCP/GPPALG
						LGPRSPAAS/C/PPLPEAPPACSP
						FCP*GEISSSGPLEPVPQV*LPGD
	}					LSEPPLGTSPLGRAPGLCSLLRA
						P/RPGAACPPASPSRPAMLLS*E
						APRRPGPSGETSPSPSLAPSEASP
						G\PPR\HHNGRDSEGAGTRTPGA
						SGSSKG/LGGQPAPHGPGGR*G
						GGRYRGSAAGSVHSAPGGCSP
						HAPGCA/CDVRG/AKPSHFHSG
						CQSG*GHQTSGGCPHLGLSPCS
					}	TCLCAGQLQQQK/PGSSSGNL*
						GQGHSSGDLGSAPPHLVPG\PG
						PLTAGGHG/SLVGDSEWGSPAP
						GAVLGQNSSGPDPQRAGRWLC
						SRAEKSGCYTAPPLG*LSAQFQ
						EEKASWVPPGSGQLEDSHPDPY
						TALPLAAPPWGTGTGCPPAWP
						RL/EPDDCGPRHLSPPTCSTLPS
						WGGCAGS
14103	44471	A	14185	264	643	RCLRPGEGRSRQGRRSRTGRGK
						AEEGGARTRQQRRAALLPERS
	•					ASQPGPRDSPRWRRRRQTTRGR
						ARPASSGGT*GSSWLTLAWPPR
						EP*RLRSASAQG/CRRQRLQQR
						RRGLYQPHRPATCGPRAAD
14104	44472	Α	14186	114	1039	QSQRLRPLADTPRRSNRPCSAA
						GLLFPQSFWSQELSQGAFLLPPS
						SPCLHCLPGTAAASRHQRSPRT
		ł				RPSVQERRETHGLCAHRCEWR
						AGSPAHGP*AHRFVSRAGSPAH
						GP*AHRCISRAGTPARGPYSRR*
						VSEAGSPA/AVVSAPCSLSS*AC
						VTG\CSPAHSP*AHRCASRAASP
						EHTSREDAEQAP*MQGLCPVH
						QEQPPSRKSVVTSPHIPPETDCR
						LLPKNPCANKHQPTATGGRAV
						PPEVQHVPPRSGMTLGSHQPLA
						LQFQMQATDAARLGLSPQGQG
						SPPGPRSTLLLGCLSLFQLSLLR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14105	44473	A	14187	1	1629	MIIYVENPKQSTSKCVRAYTFS
						KVAGHKAIEKRLEKLAVDPHR
						DRAQMLKVENVQQAWQQWIN
						KLPPARREDEDVKEIRWMIEEL
						RVSYFAQQLDHTGVDRGQPGA
					:	STWSASLMCSVTQPKLKTRKP
						GFLLGLGERNCFSVYGAGRSW
					İ	GRGLGSCGFWLPPRTRIRAADV
						CLARWGWRAHRVRVLAGKNS
						RAESMLWSFHKAAIEAARTNII
						QAQTRVEAAQATERRIAADIDD
ŀ						SELKAPRDGRVQYRVAKPGEV
					}	LAAGGRVLNMVDLSDVYMTFF
						LPTEQAGTLKLGGEARLILDAA
						PDLRIPATISFVASVAQFTPKTV
						ETSDERLKLMFRVKARIPPELL
			}			QQHLEYVKTGLPGVAWSFLYSI
						NQTICLRLDSIEAKLQALEATC
						KSLEEKLDLVTNKQHSPIQVPM
						VAGSPLRTTQMCNKVR/CVNP*
						ATVVPPPVPQPTTQQYQGLDA
l		1				GATTRVPRSLLRSEGSQTQKSP
ļ						SCGSHSQDNSGGRKAVGARRD
						RWWLGAGKGPDCRGAKCGLP
						LSGQKCPTLDPYQVVGCTSKTG
14106	44474	A	14188	3	1010	GLQTQLVPLSSPVASLDNFSNL
114100	1777	'`	14100	ľ	1010	FGADLHVEKGQVPSSDQHLSQ
						RCHFPKAPPWSVCKALLPRSNP
						ATSLSGTSKPNSGGDFNSVTKS
						QPHCELNSFVLTGHSPVLWMM
						SEHDLADVVQIAVEDLSPDHP\
						VQSCGTVVVLENHVVTDEDEP
						ALKRORLEINCODPSIKSFLYSI
						NOTICLRLDSIEAKLQALEATC
1						KSLEEKLDLVTNKQHSPIQVPM
						VAGSPLGATQTCNKVRCVVPQ
1						TTVILNNDRQN/DHCSQDGRPL
						EQQGTGFPGKMSLASQ*HGTNP
						TFPCHCAESHENKSTSLSQGTV
						QHTLHLCGHRVEPGRHRPSSLR
						SGRAGWAMLNA
14107	44475	A	14189	1	1710	OSIGIO WANDINA
14107	44476	A	14190	52	457	
14108	44477	C	14191	1	838	
14110	44478	A	14191	92	514	
14111	44479	В	14192	1	531	
14111	44480	A	14194	1628	2008	
14112	74400	/A	1777	11020	12000	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14113	44481	Α	14195	2	1616	CPPPRRPRPRAPPHHRPPAAGA
14113	14401	1	(41)3		1010	RTAAAGDRSRHGPASRVGTGP
						GQQMRKTYCRNSEPSDLDQGG
	1					GSRGGPRRQSKCVRNWWVLGL
						TDFKNEAADPHDSGAQLASPSG
						FRTRAAGGAACQSRAMRSYSS
						ALGWSMGLGAVEQGVVLVGE
	ļ					AQAAQEPMEWVGGSGMVGCR
						1
						SRAPPAGRQLRPDKKLSTAPVG
						RHRWRPSTPSAATGLGAKSLIA
				1		RGQQGRLAAPNKTDFKARAVK
						KDKEGHYIMIKGLVQKESITIINI
						YAPNTGAPKFTKQLPLDLRNKT
						DSNTIIVGYFSTPLTALNRSSRQ
						KVNKETMDLNYTLEQMDLTDI
						YRTFYPTTAEYIFYSSEHGTFSK
						TDRMIGHKTSLNKFKKIEIIPSTL
						PDHSGIKLEINSKKNLQNQTNT
		l				WKLNNLLLSDHWVNNEIKMEI
		l				LKFFKPNDNSDTTYQNLWDTA
		l				KAVLTGKFIALNAYIKKSERAQ
						VDNLRSHLKELEK*EQTKPKPS
						RRKEIMKIRAELNEIKT/QKIQKI
						NKSQSWFCEKINKIGRPLARLS
		L		100=	1.50.5	KKGREKIQISSIRKK
14114	44482	Α	14196	1097	1686	APRPLALRFGPLPSNSGAQLASP
						SGSCTGLQVELPANSCTVRWQ
						VSALGWSMGPG/GPGAGGSAD
						QGGSGGTGAHGGVGRFRHGG
		l				AAGPEPCPMGEVAKAPARNRA
						QRQWAGTAGGPCASS/NSC*PG
						C*APYCPGWWGQPAAPSAGPP
		1				SPRPPRTRAGPQAPWAAPVPAH
		1		1		ASP\PHLPAS*GSPLRPGIQGSKT
14115	44402		14107	647	1406	YQKSH
14115	44483	Α	14197	647	1496	FLNLLSWQNLMLCGLKKKRKQ
						FKDYEKKVKELNEERDKYRKG
						SVYIYTFGKEPTLNKVLDACDG
		1				KFQAIDFITPGTQYFMGCRWSC
		1		1		LPVLHCALAGLSPWVVDGTRR
						PGAVGSAGRGGSGGTGAHGGV
						GRFRHGGLQVRSPAHGEVLRPS
		Ì		1		EKSSAAQWAGTAGGPCASSTA
				1		ADLGAKPLTARGGGPAG\PPSA
						GPPSPRPPRTRAGP/PSTVGSPG/
		1				SPPTPLPPHLPAS*GSP\SGLGQP
						RKGLPQCSGGLKGWLLKHRQS
		<u> </u>	<u> </u>	<u> </u>		GCPGRGGAQSERGLRGLPARC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14116	44484	Α	14198	1	560	MAALLLVGASWL/LLLLVGAS
				İ		WLLARVLAW/TYAFYDNCHRL
İ						QCFQQ/PPKRNCF*GHLSLCGA/
				:		TEEDMRLMEDL/VPLFRDVQL
						WWLGSFYPVL/SLVHPTFTAPV
						LQASGG*PGFTTTMGLSFKRAK
			:			WQRLA/SEGSVRLEMFEHISLM
1						TLDSLQKCIFSFDSHCQMVSPG
						SCTTSRGPRIPGALPTGGARAPE
						NGDPKEIE
14117	44485	Α	14199	175	2225	RWSSVSFASGMPSAPCCWGHP
İ						QCPSAAQPTGPSWAGCRGLTM
						PNGQASPAS*VPPGAFPFLCIKQ
		İ				PLWENLMTGVGKFSREAATLE
						GPTLPTR/GVGPQCRQIADLPLE
	1					RTEGWRGD\GNPVRGVRQKRK
				ŧ		QLVAGETAASLLPCQASGESCS
						VP*QARGPPVLFLPQ\WGRRWA
						KGCASRSCF*PDFRPSHAVPGA
						REGAGLAAAPKRVARLSPAVPP
						ACC\GLQGGSSSAAQAL*PQLS
		ı				GPRLPREK*GA/PCWRGGPRCC
1			:			PQNSAPETLEYVEPRGTGAPGT
						THTGLPSPSQGCRGAEGW/WGL
						PGLPGPRAGSSPQPRGRAWRPR
						ARTPPPASWAAKGRGRPLAAG
						GLARPRDPGPGSGVAPAGSLRP
						HSSGRGWGRRRLRARPTGGNR
						GSEWPR/RPPGYPGARPGAPSVP
						RPQP/PAA*TRPLPVGCGARPFH
						PVPATV\VPHAGPP/SGMGRRRG
						RYPRPASARPAGRCGAGSAEPG
				,		RGTDKGGDGGEGKTKRGGG/L
				•		GGGAGAPKAAPGAAEPMGGR
						GAGAGGGGA\R*GGGGGGSG
						TTGSGSSSRPRPPRTRAPPETDP
						APRPRPGLVRAAGGGAASGSG
						TGRGAEGCGVRAARGVPASLR
						AALARGAQARVLVSGPAPGWA
			1			EPRPAGGARGVGVWGPVLAGP
						GRGPRGGAGRGPRALSRGCAA
L]		*AWTPGRGAGQRGPGSGRGSG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14118	44486	A	14200	<u> </u> 	2850	 MHPQERTAAATNAAIIITDGAA
14110	177400		14200		2050	TQMGVTSTPAPGHPRDGYYQG
						KDVSLLAHTHSFKISAAIEAKTT
						WVGMCRWGCKVALDVTFDVH
						QGKPEGAGRNKQIPFIQDFPFAL
						RSEKRPSLPATISGKTLPLNTKT
						CPQLLQCKREGHSLQCLIWLQ
						MVPQLGRLAVGGCHSNAWFPS
						KKAVIWEKFKHNLSGLVSDKG
						GKRTGKGLGECLLMMRKMSP
						ALEALTTFDSSLQHHFPRTFPVS
						ALEALNPGKPRQLVTPA
14119	44487	Α	14201	1	1570	MQSATSDAGSAPKKQKKVMTL
						QEKSGIA WYFFTAAQEWPNTPP
						KLRTCTESMGCCLLDCCCTGQ
						GDWTRVTLYCRYPFILFGTNA
						WVANFERPRMNANSLLASPTG
						LSPYLRFGCLSCRLFYFKLTDL
						YKKTTLGSLPDAAMAFVNCHG
						AGGTVAVRTSRGHSRCHFGFG
						VSGTRYYHLTPQVVWERQQQK
	-					VDSLLGTQQMLEAQGREF/PGP
		1				CIPLPVPTNAPAAKTTPQPPSPD
		1				L*PPNYLFLCVDGHRPIC*EPKQ
						NKSDSFTSQA*SQDG*YCGCS/C
						RAPRPHGTSATIPMSSSCKRFQP
						SSPF/CKGPYHFRFLEERPLASRP
						SIHPSCGSYWLRLGCVAVGRTG
						/SRLNLQLHSASGLQALSKQR*R
						YCQSAGAGSPEPPRGGTR*CCR
		İ				EAAPGCRAGTCPSPAANLPSLR
						RK/PGTC*PQNPPGRCGSCGASA AVLAETPARIRRSFPRARQSCPR
						TWMAAGAPGSSRRQGTKHVT*
						*SSTLEKQLDNPTATYFSVPTRK
						PVH*NQQKDL*VTG*LQIFHL
14120	44488	A	14202	111	356	VSGPPR*APWCPCLAAPTGSTS
14120	177700	^	14204		1330	HGSTARDSSTACKVS/GGATCV/
						SGCGPGLGVLGRGLSSTRSCCM
						PAG*CPSSKDTCEKSGGRTQPV
						PPFEWAAQDASLVRRCRILAHR
						PCRGTS
14121	44489	c	14204	333	473	
14141	77707		17204		17/3	<u> </u>

,	SEQ ID NO:	•	SEQ ID NO:		l .	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence		09/340,217	sequence	or peptide sequence	deterion, (-possible nacieotide insertion)
14122	44490	Α	14205	344	1035	QSAAQWFWWPGRSASLGGAK
						GMQPPSLASWPHPRSIRCLRAP
						AP*QFTNATDPRFRSAPRRSGQ
			•			QQPRGSSGPAALASPSPRQPNR
			•			AGALRTGGAGWATALDTWSC
						PRTRRAPMGSCL*AAAQQCV
		i				TGATWPPPSAIA*QSAAQWFW
		1				WPGRSASLGGAKGMQPPSLAS
						WPHPRSIRCLRAPAPCSGPSASS
						AAVQVACCCSLACCGPSRPASQ
						GHLDGTPITSPETCTTSLWSPQR
						RRAAGPPKVVDIPD*RGSRELPS
						GRGAPNTTCIPYGGSPGAKWIE
						GCPAQ/RPSPPGLHPAMRASAL
						LLMPFLQRRHRSATRLLLRPGS
				:		DCTGKK/GKGFLAGPETKRQKR
						LCPSPKTLGRALGQQEALPANV
		ļ				RC
14123	44491	A	14206	41	292	
14124	44492	С	14207	480	1637	
14125	44493	A	14208	3	410	SPPCGAPPPPVFCAPAPRAPAPP
						RAPPLAEAALPGFASAAGSAPA
		İ				RSGPAGAG*GSRSRPRARASAS
						LAPASAGDHLGYPSATAAAAP
						ASRPWLWRPSLNAARLGDPAC
						GWQASQWAAGSA*RAGSQWA
						AGEARA
14126	44494	A	14209	1	1274	FFFFHPLRSLLGDLGLHSQERSA
						PEIKPGALESTWSQARGAALRL
	į					AEAR*RAAPAAGTACSAGGAA
						SSPR*GSRATRRRRRGPTAPSRA
						ATAPGRGAAASAAPAARPART
						ALMASARGPP\TAASPAQALTR
						SAPGHLPPMSELGAVTALCESC
						PVT/AAPRSGGGEAGGTLASEW
						GRTCRNYG/PAPGIWPRPPLSAA
	ļ	}				RPAAASPPARRIWSPPAPVHSA
						ASPIGLRVAAPCSSGSAVPLRSR
						RASPPCGAPPPPVFCAPAARAP
						APPRAPPLAEAALPGFASAAAC
						RGKLRLALQGGLGRSGSRSRQE
						GPAAGS*LRTGKSWPSPPPGSA
						PLPLAQVLQALVEAGSRSRPRA
						RASASLAPASAGDHLGYPSATA
						AAAPASRPWLWRPSLNAARLG
						DPACGWQASQWAAGSA*RAGS
						QWAAGEARG
			14210		335	

NO:						Amino acid sequence (X=Unknown,
	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ļ				sequence		
14128	44496	Α	14211	344	1062	TRAEVTQQKLSRFASPPPVLRL
						RSYVSPPAAPTADPRWRPSWST
						SLRWVVKPNVSAYRIPPRASNR
						GYRASDWKLDYA*\WTGRLRIT
						SKGKTAYIKLEDKVSGELFAQA
						PVEQYPGIAVETVTDSSRYFVI\
						RISDGTGRSAFIGIGFTDRGDAF
	·					DFNVSLQDHFKWVKQESEISKE
						SQEMDARPKLDLGFKEGQTIKL
						CIGLCSKPGTTAIQLGPVLNGIG
						RTLRTDLRNKNDLEGTNL
14129	44497	Α	14212	573	1017	TDQGVGSPHCPRFQQGPPRSPR
						FQQRSPHSPDSSRG/PPHSPRFQ
						QGSPHTPDSSSG/PPYSPDYSSG
						HPTP/HTSQQGIPC*DGHGWLPC
						GPVFLWSEAPHLNPGLGVWGQ
						PRAPALPGNSAESKLWPHDRPA
						RPETPGAGTAWAHQALRV
14130	44498	Α	14213	707	836	
	44499	A	14214	724	927	SFMDFRFVLLFFHLVPNPK*RP
		-				VS/SPPGSKLLVPPWGPKT\PVPP
						HPSPPPPAHGEPGLPPPPPPPLPH
14132	44500	A	14215	29	157	
14133	44501	Α	14216	ı	363	
14134	44502	Α	14217	770	952	
14135	44503	A	14218	1053	1995	KSSEPDLSHSFIFCSPNFPPSSSL
						KRPH\FQSSILACVFCITADSSRA
						PCVLRVCDGFAGLTRSSAPSSK
						EKPCGFSLPEFTPLAVSFPQAAL
						CSLGTVSCLRPRPQLTPSWECP
į						VLLAGAVCPGLPCAAIGCPWA
						GEEGCPGATSQLDQVSWPSLQE
						APEVLWPCLDSWGWFWAPSTV
						IPLMASPRFGFMGRSQQPVLIN
				,		HLIKEPKAKFSNKNHSIRCCQG
						QRKWRRGNAAGMCCQPQIPGK
						GEVKTEFQPVSNRSVNYGCTM
						HFQETLWTPRLGLGDGDSNNN
		:				NNNNNNRITANPGKEAEKPE
						HPHLLVKM
14136	44504	Α	14219	482	833	THQGARGCTHARVSVREMRTR
				İ		RGRWATGMRVPAPCSSVHTTA
						VQRQQVADQLQTTAELSME/IL
						N/KWDQDPEAFVRKVATSYET
						WIS*YYPEDKAQSKQWLPRSGS
						GATLYSATNQYP
	l	A	14220	195	367	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14138	44506	A	14221	989	2210	LLVSCAILLSVSFLHSSSFVRSRS NRDSPACVDATYRGCSCQGEG VCGEACSCFPDLNQPQGVHAG EGVLPCSGWEDTFGKNTCLVS YQQIPNKRRPCTCEECGKAFGQ RSHLVQHTSEKLYACQECGCTF SNNSSLVKHWHVHTGEKPYMC GHCGKCFRESSSLAKHQRVHT GEKPYVC/GESTHLVQHW*FHT GEKPFAC/HR/CSKAFADFSALL ACHGTYTGERPYECRVCCKAFS PSLSLAEHIRCHTGEKLYACQE CGKAFSHSSSLSKHQQVHTECG KAFSQGSYLTQHLKIHSGEKPYI RGEHPYACGKCGKTFSHSKFLT QHEQVRMGEKPFMCGDCGRAF MQTSSLALHQRTHNGEKPYKW NECGKSCIQMSHLTEYYQNLPE KGSKDPHTDATVHSPVMD
14139	44507	A	14222	1	245	PPFRKQQAQHRNKTVLVSRSQP VPVTVPE/PSA*RSTPEGPRRAG CDPRP*PCPAGPELPAQMKVLA IEQLLSLHFKKALLF
14140	44508	A	14223	477	947	LCSLSSLPPPPPPPPPPPPPPPPPPPPPPPPPPPPPPPP
14141	44509	В	14224	438	1437	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14142	44510	A	14225	584	2866	WHFAKLSRNTQDHTRPTLGAA
						QAVPATGQPAAPGPAGAGPEW
						VPFLAVYKEQLVRNDSVLKVFF
					1	FPPIFYREGWVCCSVVESHLQE
						GFVWLFSLQL*PPKLRSLFSLSY
						CLSP\SDIEISRGQTPKAVDVLA
				į		KEIGLLADEIEIYGKILVSSAGF
						A*VVWEIL*NSLSPPELRITPTPL
						GEGKSTVTIGLVQAVSLGLCPS
						WQLYLVTPALQPGQQSETLFLK
						INK*KDILFL*FNLHLTGDIHVVI
	}					SCELSKFMDN*LLCGLIFVLCFF
					1	RGIILVTLCFLFLLKLLTFFLLNH
						*KLGINKTDPIRTMPDTHQVAG
						PYLST\DD*SVLFQ*SPAPCGNG
					1	HPTGTVADHYLCLGW/YQAQF
						DIAVASEIMAVLALTDSLGREV
		1			:	PRG*PLPG*YFVRNLMFFSHLPQ
		1				IDLGFHASVCLLGF*FAFSLL*G
						TPVFVHAGPFAADRDLCPYQFL
]	LCQTHIYKIHVIMAEYFQRYKL
						MMYAKQTSPLCDLEQAISLSGC
						VPCRHSHLKNEKFSCTGC*QYL
						PKLSLPNSDPLTLPFSADYGFKF
						ESNHANHSGDCQNLPLFLEDNS
						TYSTCFPTVTEAGFGADIGMEK
						FFNIKCRDVVLSFRSVV*AFLLL
1						AV*LGRFGGRQYTGVWVLLFS
						FLQNIQLVADGV*MVSLSLKT*
						LNGYRCKTTNIFFTRTDTRAEID
						LVCELAKRAGAFDAVPCYHWS
						VGGKGSVDLARAVREAASKRS
14143	44511	Α	14226	3	587	YPASAGLMLQNFGVIGLRYHF
					!	AIHSPAAGGLLDGLHAVAAIQG
}						ITKIETTPNHQRAPAHWTLLTQ
						QAHLQPSPLHFNLPLTLCLMHC
						PTAIPHCFADARTWVNLPTSSLI
						GHKKENLKEFISGSLIVHEILEE
						VLQAEGDFQPFTRVTVHWGKG
						NDQTFRGLLDTGSELTLIPGDP
		1			ł	KHHYGPPVKVG\AYGAQLL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	İ			sequence		
14144	44512	A	14227	2	1256	CHCGPP/VKVEAYGSQVLKGVL
	1	'	1,422,		1.200	AQVQLTVGPVGPRTHPVVIFPV
	}					PECIIGIDMLSSRQNPHTGSLTG
	-					RVWTIMVRKAKWKPLELPLPR
						KIVNQKQYHIPEGIVEISATIKD
						LKDAGVVIPTTSPFNSPIWPVQK
						TDGSWRMTVGYCKLNQVVTPI
ļ						`
						AAAVPDVVSLLEQINTPPGTWY
Į					1	AAIDLANDFFPIPVHKAHQKQF
					!	AFRWQGRQYTFTVLPQGRWEI
						NMTKIQGPSTSVKFLGVQWCG
						ACQDIPSKVKDKLLHLVPPTTK
						K/EAQCLSGFRREHIPHL\PIYRV
			ŀ			SRKAANFEWSPEQEKALQQVQ
					Ì	AAVQAAWPLGPYDPADPMVLE
						VSVADRDADWSCWQASI/GHK
						VGHAQQHSIIKWKWYIRDWAR
						ADPEGTTKGQGQRRWWQLAE
				}		RQDSRDREAAIGERQETAVGKT
						ARDGEAVCD
14145	44513	Α	14228	155	531	GNLWSVDLRPGTPLRQNFRGTI
						RQQHSRFTKNHCSQTPLLIPRQ
						TGSGVDLSKLQQTCS*GSCLVC
						TIDLANAFFSIPVHKA/HQKQFA
			ŀ			FSWQ\YTFTVLPRLTWLQPC*V
						PNLPAAETNTEPSNGT
14146	44514	Α	14229	1	518	MTVDYCKLNQVVIPIAAAVSD
						VVSLLEQINTSPGTWYAAIDLA
1						NAFFSIPVHKAQQKQFAFSWQG
1						QQYTFTVLPQWYINSPALCHNL
						IRRDLDCFSLPLDITLVHYIDDI
						MLIGSTIKWVVHSS/DSIIKWKW
1						YVHDWARAGPEGTTNGLAG*S
						GTCKKHEWKTGDKGIRGRG
14147	44515	Α	14230	281	1140	VRVLSPVEKELKLWKNTHKLL
						SYPTVGAAVTQLQNLTAMGVI
			·	1		GSHGARGQVVALNRQRQGDL
		}		1		QPFTRVTVHWGKG/NMQIFGGL
						LDTGSELTLIPGDPKHHCGPPV
						KVGAYGGQVINGVLAQVQITV
						GPQTHPVVISPVPECIIGIDILSS
						WQNPHIGSLTGIMVGKAKWKQ
						LELPLPRKIVNQKPYCIPGGTVE
						ISATIKDLKDAGVVIFTTSLFNS
						PIWPVQKTDGSWRMTVGYRRL
						NQVVTPIAAAVPDVVSLLEQIN
						TSPGTWYAAIDMANAFFSIPVH
14148	44516	В	14231	1	1521	or with the manufaction vii
14149	44517	A	14232	3	202	
14150	44518	A	14233	2	367	
L.1120_	1,1216	1.,	1- 1-3-3			

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14151	44519	Α	14234	1	553	MTVDYCKLNQVVIPIAAAVSD
						VVSLLEQINTSPGTWYAAIDLA
						NAFFSIPVHKAQQKQFAFSWQG
						QQYTFTVLPQWYINSPALCHNL
İ						IRRDLDCFSLPLDITLVHYIDDI
						MLIG\PRQLLACY/WALVETEHL
						TISHQVTMRPELPIMNWVLFDP
İ						SSHKVGCAQQHSIIKWKWYVH
			ļ			DWARAGPEGT
14152	44520	Α	14235	3	728	
14153	44521	Α	14236	1	635	
14154	44522	Α	14237	3	266	
14155	44523	Α	14238	1	1245	
14156	44524	A	14239	2	837	CHCGPP/VKVEAYGSQVLKGVL
						AQVQLTVGPVGPRTHPVVIFPV
			Ì			PECIIGIDMLSSRQNPHTGSLTG
						RVWTIMVRKAKWKPLELPLPR
						KIVNQKQYHIPEGIVEISATIKD
						LKDAGVVIPTTSPFNSPIWPVQK
					}	TDGSWRMTVGYCKLNQVVTPI
					1	AAAVPDVVSLLEQINTPPGTWY
]		1			1	AAIDLANDFFPIPVHKAHQKQF
					ļ	AFRWQGRQYTFTVLPQGRWEI
İ		1				NMTKIQGPSTSVKFLGVQWCG
		1				ACQDIPSKVKDKLLHLVPPTTK
14157	44525	<u> </u>	14240	 	533	K/EAQCLSGFRREHIPHL
14157	44526	A	14240		1043	MTVDYHKLNQVVTPIAAAVPD
14136	144320		17271	1	1043	VVSFIEQINTSPGIWYAAIDLAN
			ŀ			VFFSIPVHKAHQKQSVFCWQG
		1				QEYTFTVLPQVHINCPALCHNPI
						LRDFDHFSLPQDITLVHYIDDIM
						LIG\PV*NMLLHLVPPTTKKETQ
		1				HLEVLFGFWRQHIPHLATPSSP
	i	1		-		NGPMNKVTMMA/ANVGYAWD
						QKHGLPLTKANLVMAIGECSV/
						SPAAETNTEPSIWHHSSGQDSQ
						VQESKGGNGSGTTHHHP**STS
		1				KIFASCSHDITFCWPRGLSSRGR
		1				NAATRHNNNFIKPEVKIAFWT
						LWDPPTFKSTG*ERS*SVGWW
						D*HELPR*NQSTTPQGSVSIIPH
						GQPPHQLFGVAKARTFGIT
14159	44527	A	14242	1	336	2425.3
14160	44528	A	14243	3	1885	
	44529	В	14244	1	1423	
14161	44529	В	14244	1	1423	

NO: lo			DEQ ID ITO.	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
1	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
s	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14162	14530	A	14245	1	1361	MIISIDAEKAFDKIQQPFMLKTL
114102	1,330	1.	1 12 13	•	1301	SKLEETASPSPVVATYTPQPML
						PSAFPPLSEEINPVLPETTVMAS
						PEAVTRQDNVDSPQKPPPTPMF
						ASRPITRLKPRRAPSEEGIQRLK
						KIGMVEWISHFRPTPLSMEGPE
						HILLTNTLLNRYVKAAPASLKS
						PLTALLFMSDLTVGTTFSQLQN
						LNTMGIFGSSCDRSQVAALNHQ
						RQVPECKIVIDILNIWRIPHIGSL
						TGRVREIMVGKAKWKPLEQPP
						PRKIVIQKQYHIPGAITEISATIE
						DLKETGVVILTTSPFKSSIWPLQ
						KTDGSWRMTVNYHKLNQVET
1 1						PIAAAVPDVVSLLEQINTSPGT
1						WYAAIDLVNVFFSIPVYKAHQK
1 1						QFAFSWQGQQYTFTFLPQGYIN
				-		SPDITLVHYIDDIMLIGSSEQEV
<u> </u>						ANTLDLLPARVASWGVPYDQL
						T/GGRED*GLLHRWSCRICRHH
					1-0-	PQGSWIQES
14163	44531	A	14246	1	1795	MRKCGKPQFKLGQTNKANSRI
						QEELIHSKSLIEQEGEKPVQFSA
						FHRMWQPADSQCDIIDSADIWA
						DPLVRHREIITGSGGINRRRTRG
						GRGRGGEAAGGAENCGSREER
						ERAGVGTAVTQLQNLNTIGIIGS
		:				RGGRGQVAAINHQRQGGHSYC
						KGQQKQNSNQNSVTHVELWH
						WLINHSVPRSEIDRKPTTFLLNL
						YKQKTSRDLWPFTRVTLHRGK
						RNDQTFQGLLDTGSELMLIPED
						TKHHCGPPVKVEAYGGQVING
						VLAQIQLTVGPVGSGTHPVVIY
						PVPECIIGIGILSSWQNPHIGSLT
						SRKTDGSWRMTVHYHKLNQM
						VTPIAAAIPDVVSLLEQVNTSPG
						SWYAAIDLANAFFFIPVHKAHQ
						KQFAFSWQGQQYTFTVLPQGE
						TLVNFSLPQDITLFHYIDDIMQI
						GSSDQEVANTLDLLPRKSTTPS
						G/LYGFWRQHIS/HLGLLLTPIY*
						VTQKAA/SFEWGLEQEKALQQ
						VQATVQASLPLGVYDPADPMV
				:		IEM/SLSDPSSHKVGCAQQHSII
						KWKRYVCDQA*ASPEGTS*LY
				•		CTSFIMEKEEVC/LSLEQTLTLD
						MGLPILHAMLLWIHLWIHGLPY
						PSSWYSTQQCL

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14164	44532	A	14247	55	1241	NRFVRAAPASLKRSAGIGLLCM PDLTVGIVVTQLKNLNAMGITG SELTLIPGDPK*HFGPPVKVGAF GGEIVNKYLAQLGLTDDPVGPR THPVAIFPVPECIIGIDILSSLQNP HIGSLTGRVRAIMKGKAIMKGY YEGKGKAKWKILELPLPRKIIN QKQYRIPGGIVEMSATIEELKY AGVVIPIPTTFPLNSPIWPVRKT DGCWRMTVDYCELNQVVTPT AATVPDVVSLLEQINTSPGTWY ADIDLANAVFSIPVHKAHQKQF AFSWQGQQYTFTVLPQG\WTV NSPALGHNLIGRDLDRFSFPRDI TLLVHYVDDIMLIGSSEQEVAN ALDLLVFSDHLAIKWVMHSSIA SSSGSGICVIRLKKVLKAQPAP MASWGVLYDQLTEEEKIRA
14165	44533	В	14248	1	346	
14166	44534	Α	14249	485	717	
14167	44535	A	14250	1	345	
14168	44536	А	14251	300	689	TKRAPCSPAAGSRGRARSLALL TLYHGHYPALPPGPTPAQQPGR GLAEAAEPRGSEGGNGSNPCG RA*DGRSRREGRGRLGGWRPC CEPQPWRQAHDPAGPDRVDGG ERRGAGVLRGQDGQDEKKKPK

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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14169	44537	A	14252		2430	MAVLQPSRHRLRPQLGLEPDN WHSQQDEVSSVATARAVVPSG RGAAWMGPSGNIGGSGLVSHE CGEGTGAAGSTQLQEGMHFCR QSRWAFLTALCEVYTQSLQDT VQWLQLCQWSAQGQDDHTLIE WLMGATWTVWNDAREIPETG TGQDAWLCGSWGVDDQRPHV ELAIHCSPTNVQRVLALVDTGA DCSLVFGNPVRKPDGTWQMTV DDWELNKVTPPLHAAVPSIMD LTDHLTMELGQYHFVDYVHSP TICRGLVAMELAAWKCPKGVL LFHYIDDIMLTSDSLADLKVAV PLLWQHLAACGWAINKSKVQ WPGLSAKFLGVIWLSKTKAIPE AIIEKIQAYPRPTTMKQLQTFVG LLEYWRPFVPHLAQMIKPLYGL SKKGATWDWDNEAGTDLLAA KWAIQQTQALQVIDQGCPFELD VHVTTDGFGWVLWQRMECFR TPVGFWSQLWKGAELQYSLIQ KELAAAYAALQAYETVTDWA TVIVQMSYPIVGWTGTAQTTTL AKRGAYLEQQSMLTTTPLAAE
14170	44520		14252		220	LQEALGPIVLMEDKAMGPEAPL DPEPSLLKKGYPLVPDGAWYT DGSSQGAAAAWTAVAIQP/ITD TI*FDTGCGQSTQWAELRVVE\ MDILEACQKCPACAQAYLRQR QLPNVTQQVTVGQMPLTRWQI DYAEPLPKLQGYTYALMAADI
14170	44538	Α	14253	2	239	WWW. PORTE WAR A FORDO AND
14171	44539	A	14254	3	211	VVYYSPNEVKVVAEGFDSANG INISPDDKYGPL*NIFIFLQLWC WPNSLLCFLCQCENSFLDIYGIL
14172	44540	В	14255	13	87	
14173	44541	A	14256	42	854	VLVFLTAALFILPTFSNSMMILQ VSGGPWT/VILRRGVLLGVAPP PSLPAL\PENSVYQERQECYAFN GTQRVVDGLIYNREEYVHFDS AVGEFLAVMELGRPIGEYFNSQ KDFMERKRADQLPDCRDPLMP D*GVWISPLGQQVPNLTAQKTL QHLPSKKEPQQHHDLLVYHVT DFYPDSIQVRCFLNGQEETAGV VSTNLIRNGDWTFQILEMLEMT PQQDRT*AGVQLLEVTLNLSLF FLEAQSDSVQSKMLTGARGLFI NYLQLNLCTP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14174	44542	Α	14257	3	571	QGRATPENYLFQGRQECYAFN
						GTQRFLERYIYNREEFARFDSD
						VGEFRAVTELGRPAAEYWNSQ
		1				KDILEEKRAVPDRMCRHNYEL
						GGPMTLQRRVQPRVNVSPSKK
						GPLQHHNLLVCHVTDFYPGSIQ
ļ						VRWFLNGQEETAG\VVSTNLIL
						NGDWTFQIVVMLEMTPQ/QGE
						DVYTCQVEHTSLDSPVTVEW
14175	44543	A	14258	1	1677	
14176	44544	Α	14259	2	430	
14177	44545	A	14260	821	1428	QVTDVQDFPLKCKIQWRRTVR
						GDLQ/PFTRVTVHWGKGNDQT
						FOGLLDTGSELTLIPGDPKRHC
						GPPVKIGAYGGQIINGVLAQVQ
						LTVDAVGPWTHPVV/I/SPVPECI
						IGIDILSSWQ/NSHIGSLTGRVRA
						TMVGKAKWKPLELPLPRKIVN
1						QKQYHIPGGIAEIGATIKDLKDT
1						G\VIPTSPFNSPI*PMQKTDGSW
						RMTVDYL
14178	44546	Α	14261	507	910	
14179	44547	Α	14262	1	2898	
14180	44548	Α	14263	1	2580	
14181	44549	В	14264	159	2657	
14182	44550	Α	14265	142	377	NHLLRRQQWQTPFPPPSWS/SPR
						SSSDCCGSSENFKPVDLSLLGSV
:	;					GVGPTEPDHLDPWLQPPFQGSE
į						RFCLAGVPGATG
14183	44551	A	14266	363	638	EVWKESSGWHLAGAPLGQSFQ
						RKEQAVIFAVLQPLLVIPKQTGS
						GVDLQQTPAHLQQRGLSGCLQ
		İ				SLCSLTSS*PQRHPQNSSLVSNC
	<u></u>					VVS
14184	44552	В	14267	1	2424	
14185	44553	В	14268	446	623	
14186	44554	Α	14269	245	690	APSRSTWGSSSTAARRNLKTTR
						KLRTPRTPPHQKPRNPHHGPPR
						TPSG*TPRNPDHRPPPG\PQVGQ
						PGCWAPSQGPRGWGQLGC*RE
						EVPGGGHAGRHSTPVGG/DFFL
						QAGPWPPDASLPTNPFVLRGAS
						TPPHPLSENNKDKFLLQK
14187	44555	Α	14270	223	365	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ļ		sequence		
		<u> </u>				
14188	44556	Α	14271	1	687	MSEYIRVTEDENDEPIEIPSEDD
						GTVLLSTVTAQFPGACGLRCRN
						PVSQCMRGVRLVEGILHAPDA
						GWRNPVYVVNYPKDNKRKMV
						LGLPWKTTEQDLKEYCSTFGD
						VLMVRQSQDEPLRSRKVFVGR
						RTEDMTEDELWEFFSQYGDVM
						DVFVPKPFRAFAFVPFADDQIA
						RSLCGEDSMKGISVHISNAEPK
						RNSNRQ/LRSGRFGGNPGGFGN
					:	QGGFESYQRCERKAV
14189	44557	A	14272	1	833	
14190	44558	Α	14273	3	447	RGTAGGQPGNGPALAQSP*WC
						RLGSGGT*GKSPGTSQGRALPW
		İ				CSC/SVSTWATAAPGSQQRASP
		İ				CSWAQPGPLASLRLLMPPRLR*
						GHLGHLAWLQLFAAPCPVIHV
						AVMWSLLGIALRKVHGDWAPL
						AILSLVVCASNKLNLKKKKK
14191	44559	A	14274	988	1803	RAARVPRAPLEAVAHRWASAS
14171	77337	^	14274	700	1003	EDDGTVLLSTVTAQFPGACGLR
		l				CRNPVSQCMRGVRLVEGILHAP
						DAGWRNPVYVVNFGDVLMVR
		ŀ				QSQDEPLRSRKVFVGRRTEDM
						TEDELWEFFSQYGDVMDVFVP
						, -
						KPFRAFAFVPFADDQIARSLCG
	1					EDSMKGISVHISNAEPKRNSNR
						Q/LRSGRFGGNPGGRLQGTRYS
						PAPAPGSPGRAQGLG\AG*SRPS
						GSWLSGHLQRPEQSVGRAPAA
						PRHAVRRSLRRLAALCRRTRAR
		<u> </u>				CRRAQLHVSPQRRRH
14192	44560	A	14275	3	184	I A A A A A A A A A A A A A A A A A A A
14193	44561	Α	14276	1	352	KAAMVRFKHRYLLCELVSDDP
						RCRLSLDDRVLSSLVRDTIARV
						HGTFGAAAC\SIGFAGTIRTCQK
-		ŀ				FLIQYNRRQLLILLQNCTDEGER
						EAIQKSVTRSCLL*EEEESCEEA
		<u> </u>				AKAME
14194	44562	Α	14277	1	456	PTRAVEAAMVRFMHRYLLCEQ
		l				VCDDLRCRLSLDDRVLSSLVRD
		1				TIARVHGTFCAIACSIALALARA
						LNEEGNPEGWRDVAGASAGGR
						KKGHECPSPCTVFCTL/RYPNA
						YTVIGLLRCKKELYQVVWSVL
						YLMTYLENKGHRYPCFFNTLH
14195	44563	В	14278	34	267	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14196	44564	A	14279	3	395	SAEVGAAETTLTELRRTVQSLEI DLDSMRNLKASLENSL\GILLHL ESELAQTRAEGQRQAQEYEAL LNIKVKLEAEIATYRRLLEDGE DFNLGDALDSSNSMQTIQKTTT RRIVDGKVVSETNDTKVLRH
14197	44565	A	14280	2	1044	SSQMPSPQEGMCGKACTPALS QADSLCPLLRLASEVEGYWQM ACAEPYVQSEMNDHLSVWSLQ HGSTTLASMSFTTCSAFTNYWS PGSVQVPSYGTQPVSHAASVY AGLGGSGSRISVSHFTSF\GGLA GMGGIQNEKETMQSLRDRLAS YLDRVRGLETENWKLESKIQEH LEKNRL*LETEIEALKKELLFM KKNNEEEAKGLQAQIASSGLTV EVDAPKS/QDLAKNMADSRAQ YDKLARKNREELDKYWSQQIE ESTRVVTTQSAKVGAAEMTLT ELRHRVQSLEIDLDSTRNLKAS LENSLREVEARYALQMEPSLLF SKIIFGRVWEISYLHVFEGKQV
14198	44566	В	14281	247	857	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
٠				sequence		
14199	44567	Α	14282	1	2104	LRGSELAGGVAAPDPSSRARAP
						GRGQGSEPPRRS*GRGRAPPRA
						HTPLDDFPPGRAALSPGRGLSS
						RRPDPGSAGRPLLLCLRPPLHSP
						PPPRGCQAPAFQSLPSFHLALRL
						RSSGEEKLWGPRRAPAAP*SQR
						RKPDPARPGLAAAVPLRPV*AS
						D*GAAQVPPPSPLRDGGQPGGR
						EGVGRRTPHRAAAWLLGE*AA
						DAAAFHWDGGRPPAAPARLLP
						GAPHHREDRVHHQPRGHPLQH
						QSPGDPTPAGEQHASRH*LCRN
						PETQKLRH*TSERRDGHREEEH
						TGTAGVPRSRPATQRPHAVPAG
						GLQPHRMLPALSSGAASGGEA
						EHGQLSGRGREEDGPVWPQLP
:						AGLQGHFRGESPRWPPCLGDGS
į						EN*PGPVQAEFSGG*DPAVSESE
						DNQPRSRQFLRLQREEKAKP\D
						QRFTYLPANVPIIKTEPTDDYEP
						APTCG/RGEPGVKSSPKTILQPA
						ARDATRPQLLPRGRLPALSAEK
						HPDASSPWREPQAPRPFSRCLH
						QGRCQPGPLSPRTPAAGRRGPR
						RPG\VPRPVATHPGSPGQPPPAL
						LPQQ*MK*YEMTSPARAPTPSC
						HIGALSSAGVC*LQQTKTFE*IN
}						*THTWYHSEPPTD*MPGAEH*Y
						VQRLALQQEGKQGGRETTVSP
}						GGEVIS*QQKGGGRAEHGRPTV
						QGP/SHGNGPHAV*PHAQPFWH
						PWGSILEVPYLTRPSGHHRIEH
14200	44568	В	14283	41	4224	
14201	44569	В	14284	1	1332	
14202	44570	В	14285	140	215	
14203	44571	A	14286	234	657	VQQPGRGLDLSTDGPGGRSQV
						GLIWSCCCLH*AASGEPGGRCP
1						GS/GAPGPAGSALEFRARDGVP\
					!	GVGGPSWESHSPAAATPPPAEC
						RGPGPTPSPAPGEAAPEDREDG
						AAAPGRAEPASIVAPADGSQGQ
1						VLATQAGALGA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
14204	44572	IA	14287	134	789	PDSSQAPSRNSTRPAASTRKSSS
14204	14372	 ^`	14207		[,**	PPAARS\AGNELGSSRGDADPPP
						QNPTGPAPQSAEVPRCD/SSSA
	•					WASNGSSISPSFFSLRPWPSST/S
						KVPGAGHHTPPEGAGGASAR\S
						SSLAPSAGAPVPS\A*RSQLGTW
						LSVLRPRSL*QSRVRSPSSHVSS
						PGSMQAPNSPSSFQPGMNGAWI
						WWL*THIPLARWPGPPGILVITC
						PTVTWIKWSFICSFHPT
14205	44573	A	14288	465	1541	GPPKRKLISNFSKVSGYKINVQ
						KS*AFLYSKK/QTESQIMSELPF
		1				TIASKRIKYLGIQLTRDMKDLF
		1				KENYKPLLNEIKEDTNIWKNIP
						CSWIGRINIVKIAILPKVIYRFNA
						IPIKLPMTSFTELEKTTLKFIWN
						QKRAHIAKSILSQKNKAGGITLP
						DLNLYYKATVTKTAWYWYQN
						RDIDQWNRTEPSEIIPHIYNHLIF
						DKPDKNKQWGNDSLFNKWCW
				ŀ		ENWLSICRRLKLDPFLTPYKKIN
-						SRWIKDLNVRPKTIKTLEENLG
						NTIEDIGMGKDFMTKTPKAMK
						TKAKIEKWDLIKLKSFCTAKET
						TIRSFGSEQKSHTLVRATRLVSD
						HSQSFEAWLQFLGDTRFGTQKL
14206	44574	_	14289	1921	3199	RDSD GYLAAGQVTSRIHTGRRHFSGP
14206	44574	A	14209	1921	3199	ARCPCVSCCRRPGC*PQPAVRP
				<u> </u>		SG*HCRKTAPSHGPSIWQTR**P
				1		GSWPNLRPG*GWRWLRAQPRG
						RGIPVGCRSNLQLHPTKNPGSH
				ļ		*GSGSPWQCPGVLL*ASGRQPG
						SPSGGCT*SSGKAGLEPQWHPA
						IPGHCRNKPAAPASWERAPRPR
						AHALACGWWRYSSC*SWRS*S
						SPETPGPAQV*CHRPCPAAWW
						RCHS**TAQFL*D*FS*FLQTIGE
						PCICPKPISVPH*PMKQSH\RGIG
						GCLQYSSHPGIESSLGMSRGF/D
						RRDL*TLQIGS*RHPRAGPEGA
						AL*CLAGESSLRTPSAPCPGSPH
						TPGSSASGNTSPV**ETGCGSRQ
						GSARLVLESLLSR**CLSGPSRP
						SCLLRGISWQCGTAGRLSSSGPP
						AASGAGADAAPGGFPRLPCLEF
						GTV
14207	44575	Α	14290	3	626	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14208	44576	I A	14291	1	1065	MRGSCERSGEDEEQKEEAMVA
14200	144370		14271	1	1003	CGRLSGVPEAEQGPEANWDSD
					•	LETEGTDGLGELVRDTLYLRSC
						RAHSVVPISCFLRQGSAQELNL
						RHRGLGPQGARALASSLSSNPY
						VKRLDLRDNGLCGAGAEALAG
	1					AL/QQKQQHP*CGPVGEPAGSG
1						RSPGP/RVPPSQ/CNQAMRKMQ
						LSGNGLEEQAAQHLAELLLAH
						TDLKSLDLSYNQLNDQAGPAPS
						LRAFPQANIFLKVLDISYNGFG
						DPGASAVGEALKANNVLEELN
	1					MSNNRISAMGALSLGLGLRVN
						t i
						QTLRILVLCVRVSQVSRNPMRS EGCFGLLKSVQDNPASALELLD
						•
						FSDIQVNAEFDGLASSVRGILPG
14209	44577	A	14292	1	416	LGIKDWRLQSGV
14210	44578	A	14292	2	384	ELWVVTSFMAYGSAKDLICTH
14210	77370		172/3		304	FMDGMNELAIAYILQGVLKAL
						DYIHHMGYVHR/DLQGYDAKS
		1				DIYSVGITACELANGHVPFKDM
						PATQMLLEKLNGTVPCLLDTST
						IPAEELTMSPSRSVANSGLMTA
14211	44579	c	14294	142	474	
14212	44580	Α	14295	1	1470	
14213	44581	C	14296	91	471	
14214	44582	С	14297	195	480	
14215	44583	Α	14298	121	1130	SRSTQYELMSFLTNDASSESIAS
						FSKQEVMSSFLPEGGWYELLTV
						IGKGFEDLMTVNLARYKPTGE
		1				YVTVRRINLEACSNEMVTFLQG
						ELHVYKLFNHPNIVPYRATFIA
		}				DNELWVVTSFMAYGSAKDLIC
						THFMDGMNELAIAYILQGVLK
						ALDYIHHMGYVHRSVKASHILI
		ŀ				SVDGKVYLSGLRSNLSMISHGQ
		1				RQRVVHDFPKYSVKVLPWLSP
						EVLQQNLQGYDAKSDIYSVGIT
		1				ACELANGHVPFKDMPATQP/M
					}	VLVLSSPDAARETERHSALPVG
		}				YQHHPR*GADHEPFALSGQLW
						PE*QPDHQHPPALQR*LALPPLP
						PNLLPPLPPLC
14216	44584	Α	14299	304	412	
14217	44585	Α	14300	2	350	
14218	44586	Α	14301	191	299	HRPATAVLHVPTRSPPSPGSRPS
		1				*PPPAAALLWTPA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14219	44587	A	14302	537	5523	LARYYKFMHQDRQGVARNLL CQKEKRR*YTTKCEDNTCHWT GRTSS*PRSV/SNLNSQCGPRLW PPVTIIRQANIKTGEKAH*TSKL LPPVQLKFPARDDTLPWPGYIP* QLRRPSPR*SILSMLPRKVVPCL *QKP*TVCWKF*EPLSYIHHSLV LKMMIPMPMTLLGA*CLTQAL K*SRMERKKGRAAQRSSTLGC QSPRGSPGGGPGPPGPPGAGKQ QSNSVQTSRLLARTLI*SFSSST VKDKELARNTAP
14220 14221	44588	A	14303	2868	3095 821	PGWAGGLFGPSAGMRAWATK PSLTPPAPAGQRCPQGHRG*SG PGIYAVQPPSVPAWAAHATTQ ARLCHPRSPPLR SDRQT*RQEKRLTRRQSCCLPC
17221	1,1307					N*SFQPETTLFHGPDTFHDN
14222	44590	Α	14305	338	645	
14223	44591	В	14306	181	480	
14224	44592	Α	14307	155	543	
14225	44593	В	14308	1	1260	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
14226	44594	A	14309	1	2025	 VINIINAAQESSPMPVTEALDRV
14220	44374	^	14309	1	2023	LEILRTTELYSPQFGAKDDDPH
		1				ANDLVGGLMSDTGLEIIKDGEE
		1				EGESRTAQQYPGLSESTRQPWR
						RSRTPRTTWGWKATEQQCPDF
				:		,
	ŀ					PPPGPHPHLELLIFHCEGQGTGQ KHSTSYPCTLMSEKRGLMNVFF
						AKYTGPAMRKSSGLTPSLLHLY
						CADPLHSKPLFNCPPPTTSKTEI
1		ŀ	,			LGSCNCSPSESVRGAEKGDQEL
İ						LANIKQAEKHEKNHPEVTVAM
		ŀ				ALTDIDLQLQFSMSQPEALLLL
1		ŀ				AAGPADHLLLQLYSGHLQSAT
	1				:	GIYGEGRASSATLEDLESQYQE
		l				LAVALDSSSTIISQLTENINSLVH
						TLKEEKHEIHQVQKLGRNLFKL
			ļ			KNQTAEPLAPEPPARPSKVEQL
		l				QDETNHLRKELESVGRQLQAE
1		ļ				VENNQMLSLLNRRQEERLQEQ
						EGWQEEQERLLDKNNENKSAL
						QLEQQVKELKL\ENLEATSQ*K
		İ				QQL/TAQLSLMALPGEGDGGG
						HLDSEQEAPRPMPNIPGDLESR
						EAMVAFFNSAGANAQEEQRVC
						CQPLAHPVASSQKKPEVAAPAP
						ETGGESVCGETHQALQGAMEK
						LQESTSARGQCQRRSTGRGGHR
						QAGPGPGGDEEMLCTCTQVSS
						AGRGMGMLTVVLVVSLMMLS
			ŀ			ASFSTWCMRSISSCHCAYSAHS
						SISVFWFPREYILASGSEMMNS
14227	44595	Α	14310	3	564	SLRVSLLKAVVDRAGSRILSYIA
]				GLGVDLGFGGGSRVAGGIADL
		ļ				GRSGGGKATRSRRGSRAGGGR
						LRSRHHEEALQLRETPGPGAPE
						PMDQEYAGPGYDIRDWELRKI
						HRAAIKGDAAG/VWSAAMTRR
						FRDLDARDRKDRTVLHLTCAH
						GRVEVVTLLLSRRCQI\DICD\N*
				-		NRTPLMKAVHCQEEA
14228	44596	Α	14311	515	778	GQAQGTGDREGAVLFTARAPD
						QCRP*SCSSSRPTASSCPLGHSC
						SPGPLCTHLPLQGPPQAPTVPG
						ATATCWFSWNQCRCRAVSSYP
14229	44597	Α	14312	411	608	EAEIP*TGLPGHSRPPGPLCTHL
				1		PLQGPLGAAAWVPEAGLGQVP
						TEAAGAARAGCSAHKPWKRRK
14230	44598	A	14313	2	338	
14231	44599	Α	14314	1	390	
			<u> </u>	·	·	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14232	44600	A	14315	237	488	PWKTLDSSEAEILRGVSESSAV TLHCMEGQAVGPRQVKRIVHC *ASSA*NPHAMLLLQNR*LALC TSVPQGAAVLCGGWDPGN
14233	44601	Α	14316	358	616	
14234	44602	A	14317	13	1155	IKLPPPGCRRRGGRHRRHRYRG RAEPLAPRRRRLPPAPEQPRARP PASRFLKNTMSNGYEDHMAED CRGDIGRTNLIVNYLPQNMTQD ELRSLFSSIGEVESAKLIRDKVP GHSLGYGFVNYVTAKDAERAI NTLNG\LRLQSKTIKVSYARPSS EVIKDANLYISGLPRTMTQKDV EDMF\SRFG\RIINSRVLVDQTTG LSRGVAFIRFDKRSEAEEAITSF NGHKPPGSSEPITVKFAANPNQ NKNVALLSQLYHSPARRFGGP VHHQAQRFRFSPMGVDHMSGL SGVNVPGNASSGWCIFIYNLGQ DADEGILWQMFGPFGAVTNVK VIRDFNTNKCKGFGFVTMTNY EEAAMAIASLNGYRLGDKILQV SFKTNKSHK

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14235	44603	A	14318		2340	MFVWRNVEGHSAAVFSWYSIP FLTPPCSHTRPSNLPVTQWPPTR ENNLPSWQPLLMSVHQAQSLS ALRKEQDSSSEKDGRSPNKSDK DHIRWPMSGAHDLQQAAPGPG GAHQGHPNQDNRTVSQMLSER WYTLGPNEMQKYNLAFQNVE ARIRYVIVRGNEQGFFRMHHLR GVSSLQLGRRRPGPGTYRLEVV DMITIDARYRCSGLRSPPDPDH LKSVDLHRNEHFFLQKARWEIH LVLQATPSQIQLLHVAFVLAKQ PLRCRKQTLFLISVVGTKEKNV ASEGRNPVRPLRGHRWLPASPP AAEHFAGGADRKSRLSRAPRSA ATATPANEQRAVADVGQWAR ARGGRVRAAGRSGSCALRRSR CRRRRRHRRHRYRGRAEPLA PRRAASGQPGKLRARRGINCNA SSIPMLQVVRIKSISKHCQMAR VAGEGSLPLSHAQLTHCVFGN KQVQRTIAYLPPQQQGVQGFL KNTMSNGYEDHMAEDCRGDIG RTNLIVNYLPQNMTQDELRSLF SSIGEVESAKLIRDKVAVGLLSL CVPIRNVEIVASTSWWSLPLAP YVPRGIRRSQPSTLMDDLSCRS HQAAAPFTASVLLYISGRVKAH TAAQGFNHIPEEGVSQEAGSG WSQAWKDGWLTNVFMFSCTG HSLGYGFVNYVTAKDAERAIN TLNGLR/PPVKNH*GRPLQLGFL
14236	44604	A	14319	701	1050	LIVHFCLQVGCGVGNTVFPILQ TN/NTNSEYDPSRCFAFVHDLC DEEKSYPVPKSSLDIIILIFVLSAI VPDKMQKAINRLSRLLKPGGM MLLRDYGRYDMAQLRFKKGIL KVLNPNH
14237	44605	A	14320	701	1519	LIVHFCLQVGCGVGNTVFPILQ TNNDPGLFVYCCDFSSTAIELV QTNSEYDPSRCFAFVHDLCDEE KSYPVPKGSLDIIILIFVLSAIVP DKMQKAINRLSRLLKPGGMVL LRDYGRYDMAQLRFKKGQCLS GNFYVRGDGTRVYLCHTRCNW TRFSPTAGLEKPQKP\VDRRLQ VNRGKTTDNVPGLDSVQILQAP SVQHQLRGYLLPTRCKPVVVS GLFLKKKIVALGVV/RCL*SQPL RRLRRGGSIE/HQQSNLGKIVRD PVSESNDKIKRI

SEQ ID	1 ~	Met	SEQ ID NO:	ľ		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14238	44606	A	14321	756	1334	PGKFVLMSFLDPQPPTRILEVGC GVGNTVFPIFTNRTMTQDSLVY CCDFSSTAI\ELVQTNSEYDPSS VF\AFVQGPV**RESVTQCPRAV LDIIILIFVLSAIVPDKMQKAINR LSRLLKPGGMMLLRDYGRYD MAQLRFKKEELDTLFTTAGLEK VQNLVDRRLQVNRGKQLTMY
14239	44607	A	14322	249	474	RVWIQCKYCKPLLSSTS VYLLIVLAVLYTNNRQTESQIM SELPFTIASKRIKYLGIQL\TRDV KDLFKDNYIPLLKEI*EDTSKW KSIPCSWI
14240	44608	A	14323	1	5796	Non do Wi
14241	44609	A	14324	1	2721	
14242	44610	A	14325	1	3438	
14243	44611	A	14326	1	3306	<u> </u>
14244	44612	A	14327	1	2364	
14245	44613	Α	14328	1	2091	
14246	44614	Α	14329	1	2334	- 123
14247	44615	Α	14330	1	1986	
14248	44616	В	14331	1	2559	
14249	44617	Α	14332	1	1671	
14250	44618	Α	14333	1	1845	
14251	44619	В	14334	1	2055	
14252	44620	В	14335	1	1554	
14253	44621	A	14336	1	2559	
14254	44622	A	14337	1	1714	
14255	44623	Α	14338	1537	2204	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
-	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14256	44624	A	14339	2004	3751	YQSLAETQQQQKKNFRPISLMN
						IDAKILNKILANRIQQHIQKLIH
		1				HDQEGFIPGMQGWFNIHKSINV
		ł				IQHINRTKDKNHMIISVDAEKA
						FDKVQQHFMLKTLNKLDIDGL
						EVLARAIRQEKEIKGIQSGKEEV
						KLSLFADDMIAYLENPIVSAQN
		1				LLKLISNFNKVSGYKINVQKS\Q
		ł				AFLYTNNRQTESQIMSELP\FTI
						ASKRIKYLGI\KLTRGVKDLFRE
						NYKL\PLIKEIREDTNKWKNIPC
						SWIGRMNIMKMAIL\PKVN\YRF
						NAIPIKLPMTFFIGLGKILLLSSY
		ļ				GTKKRAHIAKTILRQKNKAGGI
		İ				TLPDFKLYYKATVAKTAWYW
						YQNRDIDQWNRTEPSEIMPHIY
		ļ				NYLIFDKPEKNKQWGKDSLFN
						KWCWENWLAICRKLKLDPFLT
						PYRKINSRWIKDLNVRPKTIKTL
						EENLGITIQDIGVGKDFMSKTPK
						AMATKAKIDKWDLIKLKSFCT
						AKETTIRVNRQPTKWENIFATY
]		SSDKGLISRIYNELKQIYKKKTN
						NPIKKWVKDMNRHFSKEDIYA
						AKKYMKKCSSSLAIREMQIKTT
						MRYHLTPVRMAIIKKSGNNRC
		<u>L</u>				WRGCGEIGTL
14257	44625	Α	14340	519	1286	LNLEERAFRHKSSEHLPHLKVH
		ł				FRHSSCGIKTKSLLHVSPPTTLIF
	:				ļ	PYPPLWGNPSN*ARPPCSVHSG
		ļ				PYPQGLASIIAHGSPGNVRGVH
						V*INDSEGGISRPSTASHKETPR
		ļ				PPTPARHADLAATRPRSLTTSPS
		1				\GAGPRA/PSRAGPSVGGLGLRP
		-	1			SGRAPRAEVALPAGLSPAGLAG
		1				PPPSGRSTGALPPAAAPTTSGP/P
1						ASTPPNAPAPGHTLSSRSPHRA
		1				ARPPPHCGAEEWTGFTMRPSW
		<u> </u>				RISQSFIVKDS
14258	44626	A_	14341	482	700	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14259	44627	Α	14342	1368	2416	SSMTQSSCPMTSYLKSKFGEPL
İ						MPSYTNPTETGVSKLVNKTQVF
						TYAHKGPVSIYFDACQAAHLSK
						LNNIWT\SVKI*DKKESASRATK
						AITEESKKECPDCDNQWTTHEF
		1				NQHLYTGRAALFASQEEKIGYT
		1				TGTCYPLNLTILKPNMTFWTKG
		ļ				HKGLLTFDQAGALLGLGIPLVI
						TKKTQRTQVQLSPIQQFRFYKS
					}	FNEHFNSEVSKIQIPPISTENLFV
						QLAKSIANNLGVTSCYVCGGA
						NMRDQWPWEARELMLQDNFT
İ		1				LPEFVTKFNANPSVWLLRNPIIG
		1				KYCIACWGKSFQNQIGETTCLG
						QQYFEESENRTQWRSFIDDSS\G
						HFNPFLQFPTLNQSWYQLE
14260	44628	В	14343	1	1071	
14261	44629	С	14344	89	283	
14262	44630	A	14345	244	315	NPKGQKDREAPLSRSVFVLKIKI
						KRAFALLLHGRF*PSTPAASASP
						ASKSPRGSGKALASALFYIVQL
	_					TK
14263	44631	Α	14346	1228	1656	QVYRPSQTPHLALSPERVAPGR
	İ					RAAGRLAPEARAPRGSP/LPPHR
						VSEKTIRVVVFHPGARRAGGTP
				-		PRAPRGDTGGAPGA\PTYSTPL
		İ				MSLHRARLESSPTGSSFPADSA
						KPVPLAVVSLDSR*GQWESRSS
						IHA\VTN*MTRH
14264	44632	Α	14347	2	705	GSWPGLDDGSATRILACPWPGL
		l				GDGPAAPKANLPLVRWKPQCF
						RGEGL*GSSCFSATLLTGLDLW
						ACCSSRSTASARLSTYPSAFTSG
						WCSSLGPTEEAPGSWINCPLAG
						ILGAATSPKPSAIPQASPEVSLD
						AEDTGMKP*DSA\GPPGPARGL
		1				CGRGCGLSPACMGLRP/PSPPA
						QSPSTALTSLT\PSPFHPPRKKW
]						SPPSAPAPSAPSPAPASLTPSPAP
L						PAPSPAPTPAPAP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14265	44633	A	14348	367	1174	WVRGQPSQSQLDLLACSTPTS*
						WPPGRAPPPPPLP/PPPPLPP\PTP
		İ				ASPPLLHPPRRAPPPPPVP\SFTP
						ASPPPDPVPRPTKALTGLP/LCP
						LPCMGR/PFVPYLGGLLPCLGP
						DFSSL**SPGGWDRPVRPLGISQ
				:		IPE\PPLTMGL*MQ*IPP*PRFQG
						HYAPLPPLQTLCISVKTPGRNPR
	1					GLPMHKILQPQNPSQPPCELPQ
						AHCPTTPPPKQCP\PAAFSILPVT
						FADKDPGAAPEGTVKAPMPTQ
						*EL*PLRPREHRVLSGPTAGQT
14266	44634	Α	14349	332	988	RRAGTPASWGHNGARAKTPSR
!	1					ACACLGLYPIHQPTARRAPAAR
		j				SHGPGPPALPRKGAAHGPRSVS
						HHPLRPCSRSSCLAWPCLHHPQ
1						PLHCHPHRHWPKWHPHPLAPH
						PLLGLHPRSRPQTPPWTSCGPSR
		l				RLPTPSGSWPAPFDRDWPN*AR
						PSALCCPFCQEPQLTPCLHLCPH
						PHPHRHLPGLSCPHRPPRWRSP
						QSPCPWWLLWWTGQWWQPGE
14267	44635	A	14350	190	718	FVLQLPPAPKSGLGRWGSPPPG
İ						QFSSSQRTGTVRGRGRCAEPRG
						SLPHPT*GNNK*HGNWMGLGS
						QGL\GRQPGEDKHCYQR/IGVG
						GAHLLRPPHQRIRSLTSTGSGCS
						HPWSRELGCSWDGQRASAPLA
						LRQAWGEGLAGRNHTIVPFHG
						DISIAPRLLPCRETLFSALCGGES
14268	44636	Α	14351	l	486	MKGEVSCKQHQDALQQVLWA
						MLQDADKQVRRAPGPVENENH
						CDFVKLREMLICTNMEDLREQT
						HTRHYELYRRCKLEEMGFTDV
1						GPENKPVRVLLFKSLSDSKRVK
					1	YDEEWIRLPVE/MKY/MLERKV
						TQKQP*VPLVFVSWRSR*ESLR
1						REEEENQQCHVLEL

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14269	44637	A	14352	1	2380	MMWSVLLQPVVLAAVERQGE ALEFLIGVPESDGENGTKLENT
						LQDIIQEDFPNLARQAHVQIQEI
						QRTPQRYSLRRATPRHIIVRFTK
						VEMKEKMLRAAREKGRVTLK
[!						GKPIRLTADLSAETLQARRDRG
						PIFNILKEKNFQPRISYPAKLSFT
		ļ				SEGEIKYFTDKQMLRDFVTTRP
						ALKELLKEALNMERNNRNFFG
ļ						VTFAFFVGGQSLGKALQETSIPF
						LREEDLEKTSILETGNGDWEAT
						RGLPEECAGMRERDCQQNEKY
Ì						QVPQFDQSTIKNIESAKGLDVW
						DSWPLQNADGTVAEYNGYHV
			:	-		VFALAGSPKDADDTSIYMFYQ
						KVGDNSIDSWKNAGRVFKDSD
İ			ļ			KFDANDPILKDQTQEWSGSATF
			1			TSDGKIRLFYTDYSGKHYGKQS
						LTTAQVNVSKSDDTLKINGVED
						HKTIFDGDGKTYQNVQQFIDEG
						NYTSGDNHTLRDPHYVEDKGH
						KYLGLIPTYDISSNIYLISKSCLW
						QANFRLNSALAQSPAITNFQFR
						GHSLKTLDLLTMKNLDSKVNII
						PVIAKADTVSKTELHKFKIKFM
						SELVSNGVQIYQFPTDDDTIAK
						VNAAMNGQLPFAVVGSMDEIK
						VGNKMVKARQYPWGVVQVEN
						ENHCDFVKLREMFICTNMEDL REQTHTRHYELYR\RLQLEEMG
						FTD\VAPKNKPASVQETYEAKR
						HEFHGERQRKEEEMKQMFVQR
14270	44638	Α	14353	105	328	TIET HOERQREEELARQWI VQR
14270	44639	A	14354	107	676	IRPDSLPSASPAQDFYAASSNTF
142/1	17037		14334	107	070	PA/DRWSALTKIYEDQTSHNIPL
						SQ/TLIQSKALTLFNSVKAEKDE
						EAAEEKFEVTRVVHIRGLIDGV
			İ			VEADIVEALQEFGLISYVMVMS
						KKRQALVEFEDVLEACNAVNY
			<u> </u>		}	TADNQIHIAGHPAFVNYSTSQKI
						SRPGNSDDPEREQCASLYHPEP
						HLFDHHGCSLHYR
14272	44640	Α	14355	221	648	LDWLRRVVGIPSLRWTCATLV
						GTRRAARHRCVNVWECGGLKE
						ITY*SVEGLCRA/EKPKPLALLE
						ETCKLPERQVVNAKEKFLEE\IK
						SATPVNT*MIRKRNCIITDMEK
						V*AVWIEDQTSHNLPLSQSLIES
						RALTLFNYVMA
14273	44641	Α	14356	122	317	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14274	44642	А	14357	338	489	WLGQN*RHGKQPGPGVGHVGT TVWS*PHSPACWAPAPQGPAPT GEEGCHF
14275	44643	A	14358	1	459	RPVWRNSQPPTAAYRWASGSR MWSTSFSRWNKTPSGKACSSS DKGPPMAQVPIPQRNVPPPTK/ QPLVLCHGEAPDFSRPPQHRPS LQKHPSTGG/PSHSPLPTSAPSSR AGLPPLAMTLDLTSGWRLWTPI QPFRLPPGSCREGTILLCSVMD
14276	44644	A	14359	609	2077	GLKAQPSSSRGHSRGPGACAAA PASRVHLPVTFCPEPNSSPCPHC TGGWGGRGQ/PACAGNPTASSP CQIECPALRSCLHALMDLGSPQ\ LRGAPRTGVPDPRLHLLPQPAL SSPPVLLRGPLLRGTCPLAARQ GQSPPAPGLLGHPGSHRGRLPP AQGGHRLPQPPGPGRRHRAPQP DPRRPGLRAAAASGPGLAAWG AVRAAERPGSGGLPGGAGPHH ARAQPGPRAAPPAAAAPGVAP AGGGAVPGGHPHRPGGEAALP AHGLPAPAAERTAPR\QPCPAC GLIARSDVHHPPVLPP/LSGPAP APGRLVGQLGVRGAPSSPVPHP ALHFPRAQP/PPGKGITTTSQTQ PAGSPPSPCMPPAATPGARHPP GSA\GHRAGV*GHTGPCDPHGP GHHGHAQLL*/DQAAVGVRGP GPWALSGMSQPPT\LPTAGHPA PACGVRAFQDGTRPPRAKPAL/ LSDKGPPMAQVPIPQRNVPPPT KHPCPGPQSSTWMRQ

1781

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14277	44645	A	14363	47	1250	EPRPSRSSGGPQLGQGLPRPMK KIPGSSAKTAGPLATRPEVPRAP
ļ						*SAGRQPWFQRPWGKRKGRKT
	1					*NHGSPGVKP/TPGPLNKDKGE
						KEERPRQQDPQRKALLHMFSG
		ĺ				KPPEKPLPNGKGSTESSDYLRA
		İ				SVTPGPWSFSP\QVASGPMPVH
						TTKKPA*APPQVL/EPKERQTGA
						AADMPQPAVRHQGREPLLVVK
						PTHSRPEGGCREVPQAA/SQNPR
		1				PAPGLQTPGTRQTS\SVTPQPCP
						PAATHSLGLGSNLSFGPGAKRP
						AQAPIQACLNFPKKPRLGPFQIP
						ESAIQGGELGAPENLQPPPAAT
İ						ELGPSTSPQMGRRTPAQVPSVD
						RQPPHSRPCLPTAQACTMSHHS
		l				AAGHDG/DPASQSALPETGKRT
						LELQPPGGPLISLS*EAGSLSRSE
						PSCVREV
14278	44646	Α	14364	23	714	EAPEIPRQNRRMTMKERPVWL
						PHK*IHPNVAKWSRVPVQVQK
				}		NPQFQE\PTPEFIPVPGTSPEAAG
						FWH*GAHDSPGTRDRSEVEPGR
İ						GHPHLGPRDSAAADPGEHSPCD
1						CP*ACPEVRRWPA\APVQIQYIIP
						SVDDFSLEFHAQDGDISDMRRE
						NVPFSPAEEGKAAPLYQQPLMI
1						PQANHMAGISPS/SPTAAEFQHR
						DVHHRGPGTRDRTHTETGRVC
				l		PHPVPTDSPAL

SEQ ID NO:	SEQ ID NO: of peptide	1	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14279	44647	A	14365	3	1869	ITTVAPSRPASSDCESDGS*EQD AGPGSRMGPEK*AGPRPTRRVG RWSQTHSQTDDTSGAVVPEYP SDR*YAGGRGPG/PGGPTVHP* KRRS*NPQETKMMARRDPKSW AKRLSPV*PRLAALLTSASPSCP LEPRVLPFRVPPSHGLCEGTSYP NSPGT*RPPCRSSAKTAGPLATR PEVPGAP*SAGRQPWFQRPWG KRKG\GKPETMEAPG*SQPGAL EQG*GREGRETKATRPAEEGSP PHVFRETSREAAAEWKRIHGVF *SSEGHPG*FPFSFPSAGGNRGT PLSCLLGVRDSTILPGCKRANA GPHNQ*EAARGPCPR*S\SAAE MSGRGSVLASLSPLRKASLSSSS SLGPKERQTGAAA/AHPSACSQ APGPRASPRGEADTQPPRG*LP RSSPGCLQNPRPAPGRQTPGTR QTSCGDL/SSPARQPPHTAWA* APISASGQEPRDLP\SSDSGLPEL PQETETGSLPDPRKRHPGR*AG GPGKSPTSASRNRTWTKYVAP DGQEDTGPGAQRRPAASAQQT LPAYCPGLHHVPSLSGQP*WGP ASQS/GSSGDWKTDAGAPASW RPPHFTLLRSREPSSLRALMCQR SLRLPVFVSHRASSMRTFRFPPP QRTAILTWS
14280	44648	В	14366	1	1519	
14281	44649	Α	14367	312	1326	TTKTGQDKDRLEEFSMQQLSGS KRKMSTQPRTEL\PPNRHTTTTS TTPATNTTCTATVPPQPQYSYH DINVYSLAGLAPHITLNPTAHP QLKQCVRQAIERAVQELVHPV VDRSIKIAMTTCEQIVRKDFAL DSEESRMRIAAHHMMRNLTAG MAMITCREPLLMSISTNLKNSF ASALR/PDGSSAIQL*SND*SCS WRPKLYDAFWDLSSLRV**PSR PEGEGRVSSEGMGESLPFSSSW PRQYQSFLCICWTDAPARNTED R*SHNKVLSSVY*NVC*NQLPC SG*AAAQSCCQSHHDPSQVLSQ PGCLCSTHCTARETLRGGHQHC HKD*SAEQGL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14282	44650	Α	14368	2	1120	PEPEPGARGGEQLNEDFEG*VA
						VPTCLHCPALSRTQSRAIQALT
						VRRGEATIRPPPCDDTKGGHRF
						LNELIKVVSPKVGAPSPSSDLPC
		ł				RMWGEERGWTVGLPEEVKIAE
						AYQMLKKQGEAHRGWGATRA
		1				CLPLPHHERPKNVIFEDEEKSK
						VRLQGGTYGDSEPRWGSVGSI
						KEMVQE/DSGRAQSTGRRRDG
						GIELGH\NVKLLTEMVMSHSQG
						GAAAGSSEDLMKVHLPPCLPHS
						LPTPQPTRTLRMRPTLFRLASDT
		l				EDNDEALGEPRAGAEVRSPPLP
		l				PPPAPVRGEEVNGDATAGSIPG
						EEVAGELGGHPSGWRGTESYP
		1				AMPTRPGEQASPEQPSASVSLL
						DDELMSLGEEGARPGGRAQQ
14283	44651	A	14369	1	432	GTRQEVTAGVEPAREWQRRLM
14203	44031		14307	1	432	EVQWGLERAPYCIQHAPR*RRS
		l				PARPGTQSCGLGACWIQ*GALS
		l				SPHCTSISRRCHSLAGSTPAVTS
						CLV
14284	44652	Α	14370	16	471	CEV
14285	44653	В	14371	66	206	
14286	44654	В	14372	234	390	
14287	44655	В	14373	207	427	
14288	44656	Α	14374	2	572	FLLERAPYGVGFSLLPGLWLGS
						RAGVGRHHDKRVGLGRAPTWS
						LEGAVSSM/TPRPQLCVP/WPC
						WGPSSPGG*QQLGQLSCSYSLV
						DR*VALLWFWMLVGSPLVKGH
						CPTPWSRFVDFSVPLATPPGIVY
					1	WGLFAVFLFSHLLPARPGEQMS
						ACCRHLHAPSSPLLTLDSLIKVY
						SCNSQGKLVRVFPPGD
14289	44657	Α	14375	175	499	KQNKSMGRKSQGFVASFGNHL
		İ				YCTASCGMPASSTSTQTWPRVL
	ļ	1				PLHSQ/ESSHGGCSWRGRR*GP
]	l				RPRPHKTPVPMVMDSTARNRA
						RRELVLAWCKSPLATALLVSW
						DK
14290	44658	Α	14376	408	625	VFGEQEVFGYMNKFSSGDF*DF
						GAPTTLSSEHCTQCVVFYPSAP
						SFPFPPVPKVHCVIPHSLVPTYE
	1	I	1	1	I	WEHTM

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14291	44659	A	14377	1263	1812	AGDAAPEDPAGRGQEDLYDEE
14271	1,,035	l' 1	1 1377	1203	10.2	LPGWGLCIPHPACLQPCHA*VT
1		1				KAG/ARVKVILPS*DTQRTKDIL
						KAGGG/PSSIHFINSAHGKPAVI
	}					VRPRGIWVLNTTPLSGQWGPYI
						GVYTAATVDMKSLGSGSLGLQ
						LALPLAASWLGVLTEAARVFG
						EQVVFGYMNSFFSGDFWDFGA
						PITRAVYTVPNV
14292	44660	A	14378	1	693	
14293	44661	A	14379	2	1299	WDYV/KRPNLRLIGVTESDGEN
11273	14001	,,	14377	-		GTKLENTLQDIILENFPNLARQA
						NIQIQEIQRTPQRYSSRRATPRHI
						IVTFAKVEMNEKMLRAAREKG
			<u> </u>			RVTHKGKPIRLTADLSAETLQA
						RREWGPIFNILKEKNFQPRISYP
						AKLSFISEGEIKSFTDKQMLRDF
						VTTRPALKELLKEALNMERKN
						RQINETENQQGYPGIELGSAPSG
						PNPLFCRHLQMSVDGRHLQNSP
						PQINRIYIFLAPHHTYSKIDHILG
						SKTLLKKCKRTEIITNYLSDHSA
						IKLELRIKKLTQNCSTTWKLNH
						LLLNDYWVHNEMKAEMKMFF
						ETNENKDTTYQNLWDTFKAVY
						RGKFLALNAHKRKQERSKMDT
	}					LTSQLKELEKQEQTHSKASRRQ
						EITKIKEELKEIETQKTLQEINES
	ŀ					KSWFFERINKIDRPLARLTKKK
						REKNQIDAIKN
14294	44662	A	14380	2	383	
14295	44663	Α	14381	1	1536	
14296	44664	Α	14382	2	1118	REHLAEGMAVGTASADLNISA
						CWLRREQQISQHSARALLRDRL
						PPQENSSWHPAGAPLGRNFQRK
						EQAAIFAVLQPPLVIPRKTGSGV
1						DLEQTPADLQKRGLTVTRKTN
						KQKAIASTSTERTPTQKPPFKSH
						QHQRPNVDKSAKMRKNQSKK
						AENSKNQNTSSPPKDHNSSPAR
						EQNWTENEFDELTEVGFRRWV
						ITNSSKLKEHVLTQCKETKKLD
			-			KRLEELLTIITSLEKNINDLMEL
						KNTAGELRDAYTSINS*IDQAEE
						RISEIGDQLNEIKCEDKMREK\R
		1				MKRNEQSLQEIWDYVKRPNLR
						LIGVPESDGENGTKLENTLQDIN
						QENFPNLARQANIQIQEIQRTPQ
						RYSSRRATPRHIIVRFTKV
<u> </u>	<u> </u>		L		L	L

SEQ ID	SEQ ID NO:	Met	ISEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14297	44665	I A	14383	1	1154	 MLPILGSRHPSSLGQARKLDYR
14237	44003	^	14303	1	11134	PSKSQCLEPGTPRACLVLYTLV
		İ				AKLIYNDYFIHYFAPRGLPPME
						KNVVFVIDISGFMFGTKMKQD
						`
						HNFSLAREQNWTENEFDKLME
l						VDFRRWVITNSSELKEHVLTQC
						KEANNLEKRLDELLTRITSVEK
1						NINDLMELKNTAGELREAYTSF
						NSCNDQAEEKISEIEDQLNEIKP
İ						NLRLIGVPESDGENGTKLENTL
						QDIIQENFSNLAKQAN\VQIQEI\
						QRTPQRY\SSR\RATPRTH/IIVRF
						TKVGNGREKMFKGQPGGKGR\
						VTLKGKPITLT\ADLSAETLQAR
						REWGPIFNILKEKNFQPRISYPA
						KLSFISEGEIKYFTDKQMLRDF\
						VTTRPALQELLKEALNMERNN
						RYQPLQNHAKM
14298	44666	В	14384	1	1554	
14299	44667	Α	14385	1	909	
14300	44668	Α	14386	2	207	DĪMSCILQNYNRPPVMALAIPIA
						VKFLHRGNKELCRNMSNYLSL
						AAITKAD\LLADHTEVIVKSILQ
14301	44669	Α	14387	496	1207	PLITGTHDSGSVESQLIKLILTM
			į			HQLFRLVLGQKDLSRAGDLF\S
	l					*NNSEIEDSLTEALEQIKIISSSSD
						YQTNNNDQAVVEICITRITTAIR
	1					ETESIEKHAKALVGLWDSCLEH
						NLRPFGKDEDTPHAKIASDIMS
						CILPELQPTPQSMALAIPIAVKF
	ļ					LHRGNKELCRNMSNYLSLAAIT
		1				KADLLADHTEVIVKSILQGMVR
		1				KLSLGTCFGRYLKVFSSSIYGL
		1				WEARPRVLEAN
14302	44670	Α	14388	219	448	VSLELCLKTTGPDCGCQTSQTP
						PAAAYHHL/PGAGLPCGPRLPA
		1				LCCSCPMGARPL*HHKR/DPSRF
						QEKTHLYLPSSH
14303	44671	A	14389	542	1365	GTEGVMANRSQEGNGGGSWR
	1	-				AKQVVVTVPF*EQSLVIMLWR
	ļ					RGGNT*QRPSEGTSRSHICSFH
						MSRHCQ*CKCLLKEP/VSAEAR
		1				DALDTKTHFPLAMIPSYLLLNG
						NIGELPEGPAGGCAQNPGLWAS
		1				/RGPARR*RSQASSSPEVRWPPG
		1				•
						QCGLRGRPCKKE\PRPHPWSRP
						CASPYRGAGGQRLRRACSAAG
		1				RKTPDRRPLPEPTRGPVSQRAV
		1				LGKLRAAAGHRIPIGDLGDGLS
		1				LGAPPAGRHLHTVLPLRREFSP
L			I		<u></u>	CEDALGIGEQQGKKRVT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14304	44672	Α	14390	3	723	PVLRGGIQEFAYIKALYERKFL
						VPKPIDYNRHAVDMELINGYPL
						C*IHHVEDPASVYDEAMELIVE
						LANHGLIHGDFNEFNLILDESD
}		1				HITMIDFPQMDSTSHPNAEWYF
						DRDVKCIIDFFMKRFSYESELFP
						TFKDIRREDTLDVEVSASGYTK
						EMQADDELLHPLGPDDKNIETK
į		ŀ				EGSEFSFSDGEVAEKAEVYGSE
						NESERNCLEESEGCY\SDHLETL
						NK*RKTVYQKRVLMHGVLK
14305	44673	Α	14391	1	1889	MGSKVKEVEDDSQMSDLVRG
						MNAFSKTGSGGVEQGRTGQGG
						TNGHIKKAKARDRDRRFKKTP
		1				QRKVGNDNLQLPLRWTEQHAE
						THIMNFCSKNYNRNIPGKPRESI
1		1				DTLNEAACRGRLSKTARKLWS
		ł				EAQDQRSFQATEMQAFLFSAK
						RIAAEKKTIIRRRDLCSIWGKLL
İ	1					PRPGSQSVPAQIPENLQEQEERA
						KWLLNGRSKFFPQTTSTLGKVT
						SNKTGVKEPELESRQPNLLKVE
						MGMKNHEIVPGSLIASIASLKH
1						GGCNKVLRELVKHKLIAWERT
						KTVQGYRLTNAGYDYLALKTL
						SSRQVVESVG\NQMGVGK\ESYI
		İ				YIVANEEGQQFALKLHRLGRTS
						FRNLKNKRDYHKHRHNVSWL
						YLSRLSAMKEFAYMKFRTLEIR
		ļ				DYFYWVKINVLTGPCSPKRGSS
İ						GEFVPCLLQLLLGFGDCWHSFG
						LWHITPVSVSMLTLLPSLLLVK
						QKKPTLDSRMGQFDFVVELQL
						YLCVLEREVIREYPPDGQIFGRY
						FDRDVKCIKDFFMKRFSYESEL
						FPTFKDIRREDTLDVEVSASGY
						TKEMQADDELLHPLGPDDKNIE
ł						TKEGSEFSFSDGEVPEKAEVYG
					i	SENESERNCLEESEGCY\SDHLE
						TLNK*RKTVYQKRVLMHGVLK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14306	44674	A	14392	150	1910	STPPGGPSVLM*STEEPGAGRS
						QPLCDDLSTRPLSCLGAGCCLW
						PAKRISAKEEPWYSSRSKDKKE
						KN\YSSPETTTSGGCHSPGDSQY
						QELAVALESSSVTINQLNENIES
						LKQQKKQVEHQLEEAKKTNNE
ŀ						IHKAQMEQLETINILTLEKADL
						KTTLYHTKRAARHFEEESKDLA
						GRLQYSLQRIQELERALCAVST
			:	}		QQQEEDRGHCLSSPDQNFSLFTI
						QSSSCREAVLHRRLQQTIKERA
						LLNAHVTQVTESLKQVQLERD
						EYAKHIKGERARWQERMWKM
						SVEARTLKEEKKRDIHRIQELER
						SLSELKNQMAEPPSLAPPAVTS
						VVEQLQDEAKHLRQEVEGLEG
1	,]	KLQSQVENNQALSLLSKEQKQ
	ļ					RLQEQEEMLREQEAQRVREQE
						RLCEQNERLREQQKMLQEQGE
					•	RLRKQEQRLRKQEERLRKEEER
	1					L/TKAGKEAVGPGGEAVEEGGE
			ŀ			ATKAGGEARALPEPQARQAAG
						RATVQLRGSEQREKERTAVGA
						ASKGAAGEARRGEGDAVHGHL
			ŀ			SAADL*EGGAAQAVTAADPVR
						GPAAAAGSLGQSGAPRSCQPPE
						PTARDPAKPRGSPWRRRWRTT
14307	44675	В	14393	463	3755	
14308	44676	В	14394	175	1835	
14309	44677	Α	14395	102	465	RAYCWAYIRLSCLTVILIAWCS
						QESALSEEEEDTTRPLETVTFKD
						VAVDLTQEEWEQMKPAQRNL
						YRDVMLENYSNLVTVGCQVTK
						PDVI\SSWSKKRSPG*WRKKCL
						GGTVQNPEEERTAV
14310	44678	Α	14396	1	330	LA LA VIDOLVED CORD LA VIDA LA CARRACTE
14311	44679	Α	14397	1	531	MLLNIPQHTRQSPRAKNYLAQ
						NVNSVKAWCSQESALSEEEED
1		1				TTRPLVRLKHLFVSCIILQCSYV
						LLPLTPEYQLNKFLYVVITMPN
						FDDFKLISIMFLQETVTFKDVA
						VDLTQEEWEQMKPAQRNLYR
						DVMLENYSNLVTVGCQVTKPD
						VI\SSWSKKRSPG*WRKKCLGG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14312	44680	A	14398	11	539	MPTTTTREMTQNSSETIKNKNG
14312	144060	^	14330	'		KKQRKQRKNSKKNRRKQKDE
]		1		DKTRRQNKRHHKKNDINPEEE
		1				RTAVRLKDGGAPAPQAVQEHD
						, ,
						AGICSASGGAQETYILVEGKGE ASTSPGQSKGSREGGVGANETC
						,
		l				SKKLYRDVMLENYSNLVT/VG
						CQVTKPDVYSSWSKKRSPG*W
14212	44601	 	1.4200	222	1227	RKKFWEALSR
14313	44681	Α	14399	322	1227	FDLKMSKCRKTPLQQLASPASF
		İ				SPDILADIFELFAKNFSYW\KPL
		1				NNEWQLPDPSEIFTCDHTEFNA
						FPWI*RTP*NEVKKPTGVIRTW
	}	1				DEWHEHTCFH**SGE/ISFPHGR
						KSVNAELWTQAWCKFHEILCS
						FPLIPQEAFQNGKLNSLHLCEAP
						GAFIASLNHYLKSHRFPCHWSW
						VANTLNPYHEANDDLMMIMD
						DRLIANTLHWWYFGPDNTGDI
		İ				MTLKFLTGLQNFISSMGTVHLV
						TANGSFDCQ*NPAGDQRMVKG
				1		HCYFPSWNWLHLSKHTQCGNE
		<u> </u>				FIFMTKKKQSLTAKSLPDTFS
14314	44682	Α	14400	3	183	KRKEITNSLDAAASRGTVPAAL
		1				AGVQYLEPRAGAAGRLGGAG
		ļ				WGRVPRPGSS/SSLSLVGR*GW
		ŀ				CPVLRTQGRCCRPPGGCWVGA
		<u> </u>				RPKAWELLSLSLVGR
14315	44683	Α	14401	21	399	IWNSAPTASRRPWSLAPGSPRS
						TWCVASTPPCAPGKAREPRDRS
						EVPPATAPAVLQSHWPCPRSSR
	İ			-		SGSAGNCVGPPHSWAPGHSQH
						LTPGPGHGGPRGP\CWHLLMQT
		<u> </u>				AWPALTPPGAMPGEAEGQ
14316	44684	Α	14402	59	478	RCSQPWHCPAALAGVQYLEPR
				-		AGAAGRLGGAGWGRVPRPREL
						LSLSLVGR*GWRPQEQDQLFPH
						LRRPVGTRSESTWGNRAGAAT
						W/RAATPSPWRT*NGTKSKQW
						TASGLWQKGDGWD*GSGKPPL
		<u> </u>				PLSVPAGQRLVPVC
14317	44685	Α	14403	1150	1319	PRRLPTACTP*RS/CSRYWMVP
						KSPRMTMMMWTKLARMGAQ
						W*PRKSNACLSRAASS
14318	44686	Α	14404	34	304	LPSRGAGLGTCSPPCLSLPPTPW
						APVRPEPPRRAPPPAPRRPVPST
						TQ/EARRSAASLLSPARPRAHRD
						GTNNSRRATLRAVTLTARVRG
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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14319	44687	A	14405	1	360	SSSLSYLPAGQGSGPAARHA*A SHPLHGLLCGPSLPNEHRPLLQ GAQSH*PPKG*GVRAHGVGLA GRSTCSP/ECGIH*VKPAGLLSFL GPRCFYEL*QSPRRSAAASLLSP ARSRAHRE
14320	44688	В	14406	80	1166	Moralite
14321	44689	C	14407	1	543	
14322	44690	В	14408	1	930	
14323	44691	В	14409	1	1473	
14324	44692	A	14410	388	606	
14325	44693	A	14411	1	798	
14326	44694	A	14412	687	832	
14327	44695	В	14413	1	1218	
14328	44696	В	14414	1	642	
14329	44697	В	14415	1	807	
14329	44698	A	14416	554	724	
14331	44699	A	14417	216	494	
14331	44700	A	14418	1	3408	
14333	44701	A	14419	330	858	VPSLDGQRGLELDTECRLVYLQ
						HSQTLC*TQGADWCIYKP*VGL QVELPASPAP\WLAFLSPWVVN GTGRHGAGGGARRGGSGCTGA HGVGGRLRHGRLQVPSPAPWE GSCSLPSPIGPPFKRAHSVLQKG SSTLMVGTCASSPKVVPFHLLS
14334	44702	В	14420	182	1513	
14335	44703	В	14421	1	1608	
14336	44704	C	14422	1	1830	<u> </u>
14337	44705	A	14423	167	433	LPSRGAGLGTCSPPCLSLPPTPW APVRSEPPRRAPPP/GSTVPSPIN HPRAEECERMARDWQAAPPAA PVRDPLGEASWAPEFFGSMLPL
14338	44706	В	14424	32	352	
14339	44707	В	14425	103	451	
14340	44708	A	14426	158	921	LPSRWAGLGTCSPPCLSLPPTP WAPVRPEPPRRAPPSAPRRPVP\ RPPKG*GVPAHGAGLAGSSTCS PDICRVCRSEGTPEKPLYHPCVC TGSIKFIHQEC\YSPDMPSRLPIQ DIFAGLVTSIGTAIRYWFHYTLV AFAWLGVVPLTAWHQLLDLGS TLNPGCSHLVILKLTTSTKTLFP NKATFILVVLSQDKTLMEDPR MEDKWKEDRFSPFNSEALEITL GERHRPQTASVIPWFCSSRHVL NLGKINFEVD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14341	44709	Α	14427	242	515	IQSVHWCIHKP*ARHRVLIGVFT
ĺ						IPELDIKVLHVPTRLRNPAGFTH
						WILHRDCRCSCMPVLRHAPALL
						SPWVVNGTGRSRAGDSTHRGG
						SG
14342	44710	Α	14428	759	988	TWPRKPLFPPRPLGL*WEGMLQ
						RSLTCPGDIFPIILVISIWLLVTY
						ANFGSWL*FLPRKWVFPFFIPVP
						NKFLIFI
14343	44711	A	14429	130	788	VLQLIKAVWTPRTQEPSWLPPV
]				NPARELQVELPASPAPCARTPQ
						LLGGRWDWAPWSRGWRSFGE
						ARAAQKPMEGVGGSGMAGCR
						SRALPHGKAAKARRAIEHSTGG
						LALLGDPVHSPQPLARVLSPSLP
						RASRAGWLLRVRGPPSPRPPGT
						PAGRQAPHAAPVPARASPSTPP/
						CRAERVGSSPGQPRKGLPQCRG
						EAEGLLKCHQSGSPGRGEELW
						HSR
14344	44712	Α	14430	108	1686	ANRCPLPHPGTGAGRCLGASA
						GRAEEAGV*GPLHEGLGYRPA
						GLFLRADTGHRTPGWGGGGG
						AGGRGGAAPGPGVGATRRFAG
						RRGCARHGAAVPAAVR\CCERL
		l				VLNVAGLRFETRARTLGRFPDT
						LLGDPARRGRFYDDARREYFFD
						RHRPSFDAVLYYYQSGGRLRRP
						AHVPLDVFLEEVAFYGLGAAA
		İ				LARLREDEGCPVPPERPLPRRAF
						ARQLWLLFEFPESSQAARVLAV VSVLVILVSIVVFCLETLPDFRD
						DRDGTGLAAAAAAGPFPAPLN
		1				GSSQMPGNPPRLPFNDPFFVVE
						TLCICWFSFELLVRLLVCPSKAI
						FFKNVMNLIDFVAILPYFVALG
						TELARQRGVGQQAMSLAILRVI
					<u> </u>	RLVRVFRIFKLSRHSKGLQILGQ
		1				TLRASMRELGLLIFFLFIGVVLF
					,	SSAVYFAEVDRVDSHFTSIPESF
		1				WWAVVTMTTVGYGDMAPVT
		1				VGGKIVGSLCAIAGVLTISLPVP
		1				VIVSNFSYFYHRETEGEEAGMF
						SHVDMQPCGPLEGKANGGLVD
14345	44713	A	14431	1	427	ST. SMQLOGI EEGIGATIAGE VE
14345	44714	В	14432	8	323	
14340	1-7-7/14	12	1.7732	<u> </u>	J-2-3	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14347	44715	A	14433	20	1050	 FNLVYMSHVSDEGALSESASAT
14347	144713	^	14433	20	1050	AASWPPHGFLQHFVPGRPQVPI
						CLYLLGQKTG*HMAPLLRQVT
						GSPIPPKTPGRGIRNMKRNEKR
						RYKTLMADGWELHQELSLGRT
						SSLPPYKGPGPWSPGFQKVSKG
						ANPVEIKRGVMLAVDAVIAEL
						KKQSKPVTKPEEIAQVATISAN
						GDKEIGNIISDAMKKCGRKGIIT
						KISSVQSIVTALEIANAYCKPLV
						IIAGDIDGEALTTLILNRLKVGL
						QVVAVKAPGFGDNRKNQLKDT
						VIATGGEVGEVTVIKDYAMLL
						KGKGNKSQIEKCVQEIIDQSDV
						TTSEYEKEKLS\EKLSDGVAVL
1 12 12	11516		14404		764	KSHPWSFCTKIYPLT
14348	44716	Α	14434	3	764	TTATVLARSIAKEGFEKISKGA
						NPVEIRRGVMLAVDAVIAELKK
:			į			QSKPVTTPEEIAQVATISANGD
						KEIGNIISDAMKKVGRKGVITV
						KDGKTLNDELEIIEGMKFDRG\
			,			YISHPFINTSKGQKCGISRDAYV
						LLSEKKIS\SIQSIVPA\LEIAHAH
						RKPLVIIAEDVDGEALSTLVLN
						RLKVGLQVVAVKA\PGFGDNR
						KNQLKDMAIATGGAVFGEEGL
		İ				TLNLEDVQPHDLGKVGEVIVTK
1.42.40	14717	ļ. —	14425	,	2000	DDAMLLKGKGDK
14349	44717	Α	14435	1	3288	MKLMETLNQCINAGHEMTKAI
						AIAQFNDDSPEARKITRRWRIG
1						EAADLVGVSSQAIRDAEKAGR
		l				LPHPDMEIRGRVEQRVGYTIEQI
						NHMRDVFGTRLRRAEDVFPPVI
						GVAAHKGGVYKTSVSVHLAQ
						DLALKGLRVLLVEGNDPQGTA
						SMYHGWVPDLHIHAEDTLLPF
						YLGEKDDVTYAIKPTCWPGLDI
						IPSCLALHRIETELMGKFDEGKL
			ł	1		PTDPHLMLRLAIETVAHDYDVI
	1	ļ.—	1440 5		100	VIDSAPNLGIGTINVVC
14350	44718	A	14436	2	422	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14351	44719	A	14437	529	2484	WGINIIKKNKKAAPRAFGVRLE ECQPATENQRWQDLNVISSLLK SFFRKLPEPLFTDEELPLAPSFCP EEENGGRGGGPSPRYRLEPMDT IFVKNVKEDGPAHRAGLRTGD RLVKVNGESVIGKTYSQVIALI QNSDDTLELSIMPKDEDILQLPA ERESGGEASEPPRVVRPEPSTRA LEPPAEDRGDEVVLRQKPPTGR KVQLTPARQMNLGFGDESPEPE ASGRGERLGRKVAPLATTEDSL ASIPFIDICVRPCAWLHALLDW MAQQRVLDMQGQSIGMGVSW DRELEEKGSEKGYDEPTSPSIDL QAKHVPASAVVSSAMNSAPVL GTSPSSPTFTFTLGRHYSQDCST SAHLNIFAGIFDISNDHLQVSTL ESNFTRIHLTHYTGGNFPAWLA
						ESNFTRIHLTHYTGGNFPAWLA WDAVPFSSDLLNLGVHVLETG AGTPFLGSACSSDGIASQMNTA GLPQIRDLPGHYYETLKFLVGH\ LKTIADHSEKNKMEPRNL\ALV FGPTLVRTSEDNMTDMVTPHA *PLTRSWETLIQALQTGSSVTKR YKGKENPCGLTRSLRQCPTLST FLPNIWQDSAPWRPGVSGPVGD LKDSTTCSSAKSKGSWAPKKEP YAREMLAISFISAVKPQAQEAA GDAGAGQQHRRRLGAGGAQS LGRGPQRRGHQERS
14352	44720	В	14438	1	1473	
14353	44721	A	14439	5	694	VPDKWGWCLFMSRYLETAIRT AACAASPRYLNPGRCSAARGPS SQSPRPPWSGHKKRTASPGWG GQSAASPMPPSRSWPASVGGPE FRLSETPSRPFTLSGMK*GWQV HSWAPALFCSVMSFSPDTCCLC K*AHSQPCSPSPSTVILLRETSSN WDWQPEDLGAPKASASPPPSTT RITRLRSSELQLRRLNPSDSCHR GFCFSFRCLLTPEHVLPKTVLRI QPGCKVA
14354	44722	A	14440	303	757	MAWKVSLRGRKVAGRAPSQLR WASPPSSGQSVKSLLLPPARRL CPGPGVPARCHRRI*GLGPGRG WRGEWRRDTNPWEPPQPRSPS/ G/PNCAWSDGCPRGAPRRPRAL *AGAARRAAPARI*VPWRGSAG GCSNRRLQVPAHE*TPTPFIRYL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14355	44724	В	14441	157	2143	RHEDHSEPTFLISPSLLCFLLPSA PPLPRLTPI\PAALLVPAVQRME PVLFLPAPGAAACRVWHPALPP PRQAAVSSSRPCY\PCVARSAPA GRVRAAGVP\PAPPSSSPRLGRV GGRSRSPNVVFRVRGAAPG/PG L*ISPWTP/PLNEAS*RQRGCPAS FAVYGGQEWRPPQWPGPPRGP
14357	44725	A	14443	258	4089	QLITCLSRPETGGKFEVDTQRP VCPPSRARAARASAAAVAAAA TLTPPTMDSFDLALLQEWDLES LCVYEPDRNALRRKERERRNQ ETQQDDGTFNSSYSLFSEPYKT NKGDELSNRIQNTLGNYDEMK DFLTDRTNQSHLVGVPKPGVPQ TPVNKIDEHFVADSRAQNQPSSI CSTTTSTPAAVPVQQSKRGTMG WQKAGHPPSDGQQRATQQGSL RTLLGDGVGRQQ/HSGQTSVQC GGGPSDPGEATCHGGQ
14358	44726 44727	A	14444	453	1326 710	SQHFGKPRQVDHLRPGV*DQP GQHGETPSLLKIQKLTGRGGAC LW/QSQLLRRLRQENRLNLGGG GCSEPGSCHCPPAWATERDSVS
14360	44728	В	14446	32	290	
14361	44729	Α	14447	244	426	
14362	44730	A	14448	69	221	QWRFCGIERGERWGKD*EIGW LPCLCGKK*IWETF\ILFCTKKN SSALGSC
14363	44731	A	14449	27	220	AVILTAKICSFTPEPSKTTSPPGG TNNSRRATLRAVTLTVKVCSFT P*GSAASFLKSVRPRTHQ
14364	44732	A	14450	115	732	EPETRIDPLSWNSSQLPVAALH LPGMSSVGAGQAICNLCCLGDL VSPSSLPALERTGPPPWNRGPK GRGRLPSL/TVSQPSLVIPPGTG KSEVNADRSRPPAYCSNLGRYS GPGPPSLVIPPGTGKSEVNADRS GPPAYCSNHFPSALP**RSSMRS LSLQQTSAWTCRHFHTS\LKSR WRFPNLNSCLLCTCRPNTTWKP CLGACTL
14365	44733	A	14451	528	875	EHGYFQGTLGQKGCKSGKPPG GTLKTLLVTLREPGRLQSP*TSL WVSERLP*YP*GDTGENCPKLS FHLRNLGPS*AG*RERPPQGGL YLPPQAYRLMQGSAFWPLFGY APSPQR

SEQ ID NO:		Met hod	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	liou	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	-			sequence		
	1	<u> </u>	1.1150	Long	10.414	LA GEOMA ENVIOLATION AND OVER A
14366	44734	Α	14452	2788	3414	ACFSYIAFWRINTTSTANLSKES
		1				LTSTTSRSWVINVNIVIFCTDWL
						FYKWLLKNFPIYFYMIFPFNQIT
						FHSMGSCAARASPTSTTPCSTA
			,			PSPIDHPRAEE\SSTRRVTGRQL
						HLQPAKICSFTPEPSKTTSPPGG TNNS\SALP*EL*HSPRRSAASL
						QFRTR*YPYRYQSQGPGSHF*L
		l		1		PLPLPSSIHDHF*LFHLQHTFQS
						HALLP
14367	44735	A	14453	662	1338	VLQLIKAVWTQRPLGGRWDW
14307	44733	^	14433	002	1556	APWSRRWCSSGRLRPHRNQWS
						GWEAQAWGLQVPEPCPAGRQL
						RPGEKSSTAPVCTSCCRTRSTA
						GPTGCPSTRRDPTLSASPS*TST
						TSLATATASASMWAGSCTSIST
						RGSASLVHQALTWPGSGPASPR
						KR\TGSHPVA*AGVRWRAHRSL
						QPQTPGLK*SSHFSLPSGWNYR
						HSNHPISQSFPHLSPLSIPQNHH
						CHHGPFSM
14368	44736	A	14454	3	415	DFLLFSEILPASQSASAILPALPA
111300	11750					SHRCPCCLDVPLPPEVPGPARLP
		1				VAAASVWRPVQHLSLLGRWA
						GPT*DRTESRETNAEVEADLPP
						LYAHGRLEPTAGPSGASAVLST
						SSEYCPGPHPKMGHRRVTSCLS
						TCQS
14369	44737	A	14455	659	818	
14370	44738	Α	14456	2	710	
14371	44739	Α	14457	349	493	SS*VRWVSGR*HPQ*STLSAPLG
						ETSGCHQGGLHDLKIVVL
14372	44740	В	14458	1	1338	
14373	44741	Α	14459	83	426	
14374	44742	Α	14460	93	345	APALCLQILLDCKSVRNRTSRF
						RPREVGIQQYKKGGGAAGKGG
						TGNAPGFLLFSKDS*KDPTPLPP
		<u> </u>		<u> </u>		PLLL*GQRDH*IQLHQE
14375	44743	Α	14461	1	1302	
14376	44744	A	14462	1	1174	
14377	44745	Α	14463	274	1428	WAL SUSSEEMED STATES OF THE
14378	44746	A	14464	411	627	HCLSHFQNTGPRGDVDFQQEL\
						FPVLMADIRKEERSHLCRSSRR
						TWTILDKS*IQGSRLSSKEQGW
14252	14747	n	14465	1	1552	GWGTSRK
14379	44747	В	14465	1226	1553 449	CLODDI*CCCLDCCLLLV*DVCD
14380	44748	A B	14466 14467	226	2061	SLQPDI*CSGLRGSLLLV*PKSD
14381	44750	В	14468	10	1099	
14382	144/30	ם	114408	110	1077	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
14383	44751	Α	14469	34	406	GGGFISLDTGSHHKSHPGGS*H
						ALGWVRPHTLNSP\PSQQPGNC
						GNPAWEC*GPHG/APSRWSCST
					,	PCRPSHPKVPAVRVAGPAGGCT
						PYRT/PTRRTPR*RPGPCAACDR
		l				P\LHPHCGPRGPGAYPR
14384	44752	A	14470	623	1671	VLRSVSSWLPTSMAMTVLAPM
						LRAMSAGRLLM*PPSMSRCPSL
						GSHSGGR*PDS/GHAGAHVPPQ
						GAIFVDSHG*GGDICGHVEVRQ
						P/QGPQSAYPGRPDGCCG*RQT
			ļ			Q*TWR*LGT*GPTQVPAPKSCS
						WPTPGPAASCGWRCAPPRTTLR
						*KSQWTCRPPGPRGPPPPLMPL
					}	VLRCASSPWPRPRLAQGQWCS
						P*APSPSGRSRCAGPRAWRKPS
		1				RRAHAAWQGCGAKRGGADSA
						GAGAPSTSCAYLATPRCAWGP
		1				HPPLGGAGLAA*GRAQPCSPAS
į						APAGRG*APGPPGPQ*GCSRPV
						AGSARSRASARRPSGRVRYGV
1						HPPAGPATLTAGTLGCYPVRRP
						PGVPG
14385	44753	Α	14471	32	972	QSGTFSTKNEGRTTRDVYDGR
						GPRVGRCGTCVYVRYVHDRPH
						APPHERARRNDGNTQRTGNAR
						PGTAMPTATTDQPKGRSTRHK
						ETKRGRTARNGKQQNNPRK/RP
						RKDKRKQGSAGEGERGAGGKE
						KGQPRDTEAKGDRRTDFPPPNG
						\P*GQSKRGANIGGWFPPPRLGT
						PNPPLPGPQSAQVSGP*TPQMW
						MPTGTGCCGGSQ\SGSPPPWCV
						SP*PVAEALFHSST/PAPRHPSPP
						SSSPKHSHQPSTPPP*ISLGQGPC
						RPPKRRPAPDQPQVPRCPEPSC
						AYLATPRPTMRMQSACRGSAR SRASARRPSG
14386	44754	Α	14472	1	3128	
14387	44755	В	14473	239	5474	
14388	44756	Α	14474	542	781	PASIIMTGSISHITILTLRVNRLN
						ASIKRHRLTDWQIR/IKSQDLSV
						CYIQETHFTCKDPHKLKIKGWR
L _						KIYQVN*KQKKK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
:				sequence		
14389	44757	A	14475	724	1340	QLTLQGRRVAPR*PSRVALEQE/
14365	177/3/	1	14473	72.4	1510	CGYNALLRYEGFENDSGLDFW
]						CNICGSDIHPVGWCAASGKPLV
						PPRTIQHKYTNWKAFLVKRLTG
						_
						AKTLPPDFSQKVSESMQYPFKP CMRVEVVDKRHLCRTRVAVVE
		•			}	
						SVIGGRLRLVYEESEDRTDDFW
						CHMHSPLIHHIGWSRSIGHRFK
						RSDITKKQDGHFDTPPHLFAKV
1.4220	14750	┦	14476	1000	2226	RSFYKNPHL
14390	44758	Α	14476	1822	2226	KRLFTNRTRITPFVAINSHHHHH
						HHITIINTIITTTSTINSHHHHHHY
		1	:			ҮНННННННННОНТТТТОТТ
						TNTTTTNNNTTTTITITTTTNTT
		1				TTSTINSHHHHHNQHCHHHHH
						нннн/нннниочнинн+нүо
						\ннннннүүнннннннннү/
						нннниднниодни
						ДИНИНУННИОННИОНУО
						ннннионсинниннини
		<u> </u>				ННОРРНННОН*НЅНОНН
14391	44759	Α	14477	232	625	EGCCCIAIMVQKYQSPVRVYK
						YPFELIMALRK\RYPRCCVTCS*
						TGGAQQVGFRFLDLPLSLCTP*
		l	1			CLWRTSSSNF*RKAAGCSGSRL
		ļ	:			QSHHFGLIPQPTAPKPWSSPLHI
		<u> </u>				TKPQTLGQRDPGPRTRKRSK
14392	44760	A	14478	1	808	MPKTTACNVHTVTWASCENAD
		İ				SDSTGLRHHDSCEDPASPGRSD
						LCLVPHPTRPAVRGGRPAGGRC
						PAVPAEGPWRHLQQAQEGLKR
						LAGGLERLLENPVRRALPQGA
						QGHRAEPTEARPHPAPEHRRAG
						AAAERAERLESQPVGASAAAA
						SAASARIPYLVAVPAAGPPRHR
						CRRHRRGPGTSSGPRTALIRPAT
	1					ARARAGIVSEPRTAPGATRARP
						WRQGSHNPVAPSAP*LLRGPSIP
						WQIRPMPCPTSHQTCSERRPSG
						RWQMPGSACRRSLETSSAGAG
]						RAQEAGGRLGETSGESG/RAGL
						YLKELRAIVLNQQRLVRTQRQS
						IDELERRLNELSA*NRSLWEHP
]						QLPQAQPPPGFLTS*PSPLPALL
						GTAAAATAGAQEHLQDHGQRS
						SARPLPAPELASFQNHEQLLAQ
						PEPDPGGRAHTTQSPHQHPGTL
						GVKANKEKKGPPGCCAAPGTP
		L	1	I		LQQESPAPSARES

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	t .			sequence		
14202	144761	1	14479	1	831	I MIRYFGDYWTQALLTLIPGDPK
14393	44761	Α	14479	1	031	•
						HPRGPPVKVGVYGGQGYINSL
			,			ALCHNFIWRELNRFSLLQDVPL
						VRYIDNIMLIGSIKFLGVQWCG
		l				ACRDIPSKDPADPMVLEVSVAD
						RDAVWSIWQALIDESQQRPLGF
						WSKSLPSSADNYSPFERQLLAY
						YWALVETERSTMGHQVTMLPE
	1					LPVMNWVLSDPSSHK/ANGLA
ŀ						GWSGTGKKHDWKIGDKEIWRR
						GMWMDLSEWSK/D/VKIFVSHV
						SAHQRVTSAEEEFNNQVDRMT
						RSMDTTQPLYPTTPVIAQWAHE
14394	44762	В	14480	56	3476	
14395	44763	Α	14481	3	591	DPADPMVLEVSEADRDA\VPIS
		-				ESQQRPLGFWSKALPSSANNYS
						FFKRQLLACYWVLVEIEHLTM
						GHQVTMRPELPIINCVLSDPCSH
						KVGHAQQHSIIKWRWYIHDWA
						EGTSKLHEEVAQIPMVSTPSLP
		ŀ				QPAPMASWEVPYDQLTEEEKT
						RAWFTDGSARHAGATQKWTA
						VALQPLSGTSLQDSSEEKSSQW
14396	44764	A	14482	1	1275	MEKNDIDQSRRRKVVEMNENV
1.000	,	'				QLVRYGEQASDLKTATKFKSPS
						EVKHIRFIQQAFIGSKLLMRPW
						TTVTKQILSLKSAKIALSESLIPC
	1					RKRAEFVEKQTQAFIMQVSDL
						QQKVHAQPSQVSTVKVKALIG
						KEWDPATWNGDVWEGPDEAG
		1				DTEFVNSDEAFLPEATAFPSPEV
						GNDOTVWGLLDTGSELMLIPG
						DPKYYRGPPVKVGAYGGQVIN
	1					RGLAQIQLSMDPVGHRTRPVVS
						SPLPECIIGTDILSSRQNPHIGSLT
1						DITLVHYIDDIMLIGSSEQEVTIT
		1				LDLLVQAVAQAALPVGPYDPA
	1					DPMELKVSVADRDAVWSLWQ
						APIVTMRPERPIMNWVLSDPSS
						HKVGCAQQHSIIKWKWYIRDW
						ARSGPEGTNHSRRVCS/LPQVQ
						APSAR*ATARRGGQAASGCCLF
	1	1			150	WSCTCFSNFDV
14397	44765	A	14483	604	1760	NSWCRWFNFRY*YPTFYWES*
		1				NT*IPLREWFG*NHDASCP*TLD
		1				SKSFSFETRTGSPCSSSLQTAYC
						GTLWIVQGV
14398	44766	Α	14484	2710	3060	
14399	44767	A	14485	3	227	
14400	44768	A	14486	3	266	
14401	44769	Α	14487	2	564	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14402	44770	A	14488	2	503	FFFFLPIGFWRQHIPRLSVLLQHI Y*VT*NAASFEWGPEQEKAL*Q VQAVVQVALPLGPYDPADPML LEVSVADKDAGCTRL\QAPIGE SQWRSLGFRSKALRSSAENYSS FERQLLACYWALVETECLTLG HQVTI*PELPIMNWVLSDPSSHK VGHEQQHSIIKWK
14403	44771	Α	14489	2	469	
14404	44772	A	14490		633	MTVDYRKFNQVVTPMAA/AVP DAVSLLEQINTFPGTWYAAIDL ANAFFSIPVHEAHQKQFAF\LPQ GYINFPALCHNLIRRELDFFLLL QDITLVHYIDDILLIGSSEQEVV NTLDLLIHKRSKEAEHTAASRIR VSCLPEQKSHEQTLPWEQVPSS GDIKEYFPNAFVLLTTASLQGQ DNTSQLQLTWKAPEDIKMSKT DADADEEIEALRG
14405	44773	Α	14491		518	MTVDYCKLNQVVIPIAAAVSD VVSLLEQINTSPGTWYAAIDLA NAFFSIPVHKAQQKQFAFSWQG QQYTFTVLPQWYINSPALCHNL IRRDLDCFSLPLDITLVHYIDDI MLIGSTIKWVVHSS/DSIIKWKW YVHDWARAGPEGTTNGLAG*S GTCKKHEWKTGDKGIRGRG
14406	44774	С	14492	33	932	
14407	44775	В	14493	27	2457	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14408	44776	A	14494	332	2158	RGYVFCSWKKTDGS/WRMTVD
		Ì				YCKLNQVVIPIAAAVSDVVSLL
						EQINTSPGTWYAAIDLANAFFSI
						PVHKAQQKQFAFSWQGQQYTF
						TVLPQWYINSPALCHNLIRRDL
						DCFSLPLDITLVHYIDDIMLIG\P
1						RQLLACY/WALVETEHLTISHQ
						VTMRPELPIMNWVLFDPSSHKV
						GCAQQHSIIKWKWYVHDWAR
						AGPEGTTTPVISQWPHEQCGHG
						GRDGGYAWAQQCRLPLTKAD
						LNTATAKRPICQQQRPTLSPQY
						GTIPQGDQPATWWWVDYMGS
						LPSWKGQRFVLTGIDTYSGYGF
						AYPACNASAKTAICGLTECLIH
						HHDIPHSIASDQGTHFMAKEVR
; 						QWAHDHGIHWSYHVSHHPEA
						AGLIEWWNGLLKSQLQCQLGD
		İ	İ			NTWQGWGKVLQKVVYALNQH
						PIYGTVSPIAKIHRSRNQGLEVA
						PLTITPRDPLAKFLLPFPATLQS
						AGLEVLVPEEGTLPPGDTMIPL
						NWKLRLPPRHFGLLLPLNQQA
						KKGVTVLAGVTDLDYKDEITL
						LLHNGGKEEYAWNTGDPLGLL
		1				LILPCPMIKVNGKLQQLKPEAL
		1				VPKGVVFPPGDTTMLSLSWKL
						RLPSGHVGLLMPLSQQVQKGV
14409	44777	Α	14495	1	3187	MAEGKEEQVLSYTDGSRQREN
		;] [EEDAKAETPDKTIRSHETYSLPR
						EWYEGNRPHDSITSQWVPPTTR
						GNYGSTIQDEIWVGDHSGYVRP
						VPVPRSLNSDISYFGVGGKQAV
]				FFVGQSARMISKPADSQDVHEL
						VLSKEDFEKKEKNKEAIYSGYI
						RNRKDDYDNHTGIDLVGTIIATI
]				KGSNEEDTDTPLFIGKVRTLEFP
						FVNGSAEIMLMPSNQQHKTDE
						KGRANLGVFSVFAPRGEHTLQ
	L	<u> </u>			167	VKAIYNKSIIEGP
14410	44778	A_	14496	266	467	
14411	44779	В	14497	152	1245	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14412	44780	A	14498] 1	1445	MLRAAAGGQGLAAAGMGLCA
						RPAGGSQLPRERMKGNRACLP
						LPPAVALTKGREKPDAGHQGR
			ŀ			STQMMRESERGPCTHWTEERP
		İ			·	GRQLKILLQKQMRQNTLMAFQ
						VSWDGRNKEIGRTFPHATAPHT
						QQLIFAMKIKRTGVTVEPWGLF
						ETGQRIRPIQAAGL*LSSSSTTC
		ŀ				VALAPLLTMHVKSLALLSAP/Q
						MGALSTITSITTTRRQTPAESHF
		1				PAARFHLLAPHSPTLPPCCPGEP
						SESAVTVLCFAKGKTAQDAAV\
					,	PTTSLTAAGHLAPRS*SSWSRM
		1				PVKVRRAPDSLPETSAHALPLT
						H*SWLPSASWGPASRALPGSEW
į						VLKPVWQ/AITGPVSQITSSPGIP
						KP\GPLQA*GSAHPQALGFSVL
İ		1				QIPLSNSPTSPMTPPLPRPKISNS
						SLPSQ/PPQALFPNRWTNSSSPV
		İ				HVITSAAQ\PRTPREWPAVSFYR
						TTATTASLIPHLWAPPQEN*QTS
						PAQNGSTLPCMSQATFHFPSP
14413	44781	A	14499	44	455	HLRNRTR/PSQITPHIYNHLIFDK
						/PLFNIWWWENWLAICRKLKLD
				*		PFLTPYTKINSRWIKD*NIRPKSI
						KNLEENLGNTIQHLG/TGKDFM
						TKTPKAMATKAKIDIWDLIKLK
	ĺ					SFCTAKETIIRVNRLPT*WEKIF
						AIYP
14414	44782	A	14500	1	346	
14415	44783	A	14501	1926	2448	DFIAVITTRKQLKTKYNAHHSK
1						QAITKHFSPKISDLILCLKKKIM
						NRHFSKEDIYAAKKHMKKCSP
						SLAIREMQIKTTMRYHLTPVRM
			1			AIIKKSGNNRCRRGCGEIGTLLH
				}		CWWDCKLVQPLCKSMWRFLR
						DL\DPAIPLLGIYPKDYKSCCYK
						DTCTRMFIAALFTIAKTWNQP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14416	44784	A	14502	1	2142	MIILIDAEKAFDKIQQPFMLKTL
			- 10 0 -			NKLGIDGTYLKITRAIYDKPTA
						NIILNGQKLEAFPLKTGTRQGCP
				,		LSPLLFNIVLEVLAQAIRQEKEI
						KGIQLGKEEVKLSLFADDMILY
						LENPIVSAQKLLKLISNVSKVSG
						YKINVQKSQAFLYTNNRQTESQ
						IMSEFPFTIATKRIKYLGIQLTRD
						VKDLFKKYKPLLNKIKEDTNK
						WKNIACSWIGRINIMKMAFPR
						WELNNENTWTQEGEHHTLGPV
ļ						VGWGKRGGIALVDIPNVNDKL
						MVLEVLARAIRQKKEIKGIQLG
						KEEVKLSLFADDMIVYLENSIV
						SAQNLKLISNFSKVSGYKINVQ
:						KSQAFLYTNNRQTESQIMSEFP
						FTIATKRIKYLGIQLTRDVKDLF
ŀ						KENYKPLLKEIREDTNKWKNIP
-						CSRIGRINIMKMAILPKVIYRFN
						DIPIKLPMTFFTELEKTTLKFIW
						NQKRACIAKTILSKKNIAGGITL
						PDFKLYYKATVTKTAWYWYQ
		:				NRDIDQWNRTEASEVTSHIYNH
ŀ						LIFYKPDKNKKWGNDSLFNKW
1						CWENWLAICRKLKLDPFLTPYT
	:					KIHSRWIKDLNVRPKTIKTLEEN
						LGNTIQDIGMGKDFMTKTPKA
	_ \					MATKAKVDKWDVIKLKSFCTA
						KETTIRVSRQPTEWEKIFAIYPS
						DKGLISRIYKELKQIYRKK\TNN
						PIKKWAKNMNRHFSKEDIYAA
<u></u>						NRQMKKCSSSLVIREMQIKTTM
14417	44785	Α	14503	209	3816	QGRPTFRFRKYREHHKDTPREE
				:		QLQDT*SSDSPKLK*RKKC*GQ
İ						PERKVKLPTKGSPSD*KRISRQ/
ŀ						KTLQARRQSWFFEKINKIDRPQ
					:	ARLIKKKREKNQIDTIKNDKGD
						ITTDPTEIQITIREYYKHLYANK
						LENLEEMDKFLDTYTLPRLNQE
						EVESVNRPITGSEIEAITNSLPTK
		}				KSPGPDGFTAEFYQRYKEELVP
		1				FLLKLFQPIEKEGILPNSFYEASII
						LIPKPGRDTTKKGNFRPISLMNI
		<u> </u>		<u></u>		DAKIL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	F	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14418	44786	IA	14504	1	2877	MENDFDELREEGFRQSNYSELR
14410	7.00	1	17307	1	2077	EDIQTKGKEVENFEKNLEECITR
						ITNTEKCLKELMELKTKAQELR
						EEWRSLRSRCDQLEERVSAME
		1				DEMNEMKGEGKFREKRIKRNE
						QSLQEIWDYVKRPNLCLIGVPE
ľ						RDGQNGTKLENTLQDVIQENFP
Į.						NLARQANVQIQEIQRTPQRYSS
						RRATPRHIIVRFTKVEMKEKML
						ł
						RAAREKEIQTTISEYYKHLYTN
						KLENLEEMDKFLDTYTLPTLNQ
14410	44707	-	14505	1	22.47	EEVESLNRPITGAE
14419	44787	Α	14505	1	2347	MELKTKARELHDECTSLSSRFD
				ĺ		QLEERVSVMEDEMNEMNLPTK
						KSPGPDGFTAEFYQRYKEELVP
						FLLKLFQSIEKEGILPNSFYEPSII
						LIAKPGRDTTKKENFRPISLMNI
						NAKILNKMLANQIQQHIKKLIH
						HDQVGFIPGMQGWFNIRKSINV
						IQHINRTKDKNHMIISIDAEKAF
						DKIQQHFMLKTLNKLVLEVLA
						RAIRQEKEIKGIQLGKEEVKVSL
						FADDMIVYLENPTVSAQNLLKL
						IGNFSKVSGYKINVQKSQAFLY
						TNNRQTERQIMSELPFTIASKRI
						KYLGIQLTRDVKDLFKENNKPL
						LKEVKEDTNEWKNIPCSWVGRI
						NIVKMAILPKVIYRFNAIPIKLP
	İ					MTFFTELEKTTLKFIWNQKRAC
			<u> </u>			IAKSIFSQKNKAGGITLPDFKLY
						YKATVTKTAWYWYQNRDIAQ
						WNRTEPSEIMLHIYNYLIFDKPE
						KNKQWGKDSLFNKWCWENWL
						AICRKVKLDPFLTPYTKMNSR
						WIKDLNVRPKTIKTLEENLGITI
						QDIGVGKDFMSKTPKAMATKA
						KIDKWDLIKLKSFCTAKETTIRV
						NRQPTTWEKIFATYSSDKGLISR
						IYNELKQIYKKKTNNPIKKWAK
						DVNRHFSKEDIYAAKKHMKKC
1	ļ					SSSLAIREMQIKTTMRYHLTPV
						RMAIIKKSGNNRKIQ/GGIWCD
						RIL*R*TTCRVAKEIQSL*RRI/W
<u></u>						KRLQRTLSIPVLDAV*PPMF*AS
14420	44788	В	14506	1	2265	
14421	44789	Α	14507	1	2832	
14422	44790	A	14508	1	2109	
14423	44791	A	14509	1	2757	
14424	44792	Α	14510	1	2304	

	of peptide sequence 44793	A	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion) MGDFNTPLSTLDRSTRQKVNK
	·	Α				
14425	44793	A	14511	1	2685	MGDFNTPLSTLDRSTROKVNK
14425	44793	A	14511	1	2685	MGDFNTPLSTLDRSTROKVNK
				i		DTQELNSAPHQADLIDIYRTLH
			l			PKSTEYTFFSAPHHTYSKTDHIL
						GSKALLSECKRTEHTNYLSDDS
						AIKLELRIKNLTQNRSTTWKLN
						NLLLDDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFVALNAHKRKQGRSKIDT
		l				LTSQLKELEKQEQTHSKASRRQ
		l				EITKIRAELKEIETQKTVQKINES
						RSWFFERINKIDRQLARLIKKKR
		İ				EKNLIDAIKNDKGDITTDPTEIQ
						TTIREYYKHLYANKLENLEEM
						DKFLDTYTLPRLNQEEVESLNR
						PITGSEIVAIINSLTTKKSPGPDG
						FTAEFYQRAIRQEKEIKGIQLGK
						EEVKLSLFADDMIVYLENPIVS
		1				AQKLISNFSKVSGYKINVQKSQ
						AFLYTNNRQTESQIMSELPFTIA
	:					SKRIKYLGIQLTRDVKDLFKEN
						YKPLLKEIKEDTNKWKNIPCSW
		l				VGRINIVKMAILPKVIYRFNAIPI
		l				KLPMTFFTELKKTTLNFIWNQK
		1				\RAHIAKS/VLSQKNKAGGITLP
						DFKLYYKATVTKTAWYWYQN
						RDTDQWNRTEPSEIMPRIYNYL
						IFDKPEKNKQWGKDSLFNKWC
						WKNWLAICRKLKLDPFLTPYT
						KINSRWIKDLNIRPKTIKTLEEN
						LGITIQDIGMGKDFMSKTPKAM
	1					ATKAKIDKWDLIKLKSFCTAKE
	1			1		
14426		<u> </u>	i			TTNRVNRQPTKWEKIFATYSSD

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14427	44795	A	14513		1544	HINRAKDKNHMIISIDAEKAFD KIQQHFMLKTLNKLGIDGTYFR KSINVIQHINRAKDKNHMIISID AEKAFDKIQQHFMLKTLNKLGII DGTYFRKSIN/DNPAYKQSQRQ KP\LIISIDAEKAFDKIQQPFMLK TLNKLDNIIVYLENPIVSAPNLL KLISNFSKVSGYKINVQKSQAF LYTNNSQTESQIMSELPFTIASK RIKYLGIQLTRDVKDLFKENYK PLLKEIKEDTKKWKNIPCSWVG RINIVKMAILRKVIYRFNAIPIKL PMTFFTELEKTTLKFIWNQKRA PVAKAILSQKNKAGGITLPDFK LYYKATVTKTAWYWYQNRDI GQWNRTEPSEITPHVYNYLIFD KPEKTKQWGKDSLFNKWCWE NWLAICRKLKLDPFLTPYTKIN SRWIKDLNVRPKTIKTLEENLGI TIQDIGMGKDFMSKTPKAMAT KDKIDKWDLIKLKSFCTAKETT IRVNRQPTEWEKTFATYSSDKG LISRIYNELKLIYKKKTNNPIKK WAKDMNRHFSKEDIYAAKKH MKKCSSSLAIREMQIKTTVRYH
14429	11706	D	14514	110	2153	LTPVRMAIIKKSGNN
14428	44796	A	14514	1	3210	MVKGSIQQEELTILNIYAPNTG ALRFIKQVLRDLQRDLDSHTIIM GDFHTPLSTLDRSTRQKVNKDI QELNSALHQEDLIDIYRTLHPKS TEYTFFSAPHHTYSKIDHIVGSK ALLSKCKRTEIITNCLSDHSAIK LELRIKNLTQNRSTTWKLNNLL LNDYWVHNEMKAEIKMFFETN ENKDTTYQNLWDTFKAVCRGK FIALNAHKRKQERSKIDTLTSQL KELEKQEQTHSKASRRQEITKIR AELKEIETQ
14430	44798	A	14516	3	2820	ENKDTTYQNL\WDAFKAVCRG KFIALNAHKRKQERSKIDTLTS QLKELEKQEQTHSKASRRQEIT KIRAELKEIETQKSLQKINESRS WFFERINNIDRPLARLIKKKREK NQIDTIKNDKGDITTDPTEIQTTI REYYKHLYANKLKNLEEMDKF LDTYTLPRLNQEEVESLNRPITG SEIVTIINSLPTKKSPGPDGFTAE FYQRYKEELVPFLLKLFQSIEKE GILPNSFYEASIILIPKPGRDTTK KENFR

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14431	44799	Α	14517	687	2274	LRDCKRINQHPVKTDQSAFCK
						M/API/MQDVVLEVLARAIRQEK
						EIKGIQLGKEEVKLSLFADDMI
İ				9		VYLENPIVSAQNLLKLISNFSKV
						SGYKINAQKSQAFLYTNNRQTE
					ŀ	SQIMSELPFTIASKRIKYLGIQLT
						RDVKDLFKENYKPLLKEITEDT
						NKWKNIPCSWVGRINIVKMAIL
						PKVIYRFNAIPIKLPMTFFTELE
						KTTLKFIWNQKRARMAKSILSQ
						KNKAGGITLPDFKLYYKATVT
						KTAWYWYQNRDVDQWNRTEP
						SEITPHIYNYLIFDKPDKNKQW
						GKDSLFNKWGWENWLAICRKL
			 -			KLDPFLTPYTRINSRWIKDLNV
						RPKTIKTLEENLDITIQDIGMGK
						DFMSKTPKAMATKAKIDKWDL
j						IKLKSFCIAKETTIRVNRQPTKW
						EKIFATYSPDKGLISRIYNELKQI
						YKKKTNNPIKKWAKDMNRHFS
		1				KEDIYAAKKHMKRCSSSLAIRE
						IQIKTTMRYHLTPVRMAIIKKSG
						NN/R/CF*WQKPGPSG*ALRSWK
						EDIAGTSYVGMNQCQMRKNTK
		<u> </u>				TLI
14432	44800	В	14518	1	3192	
14433	44801	Α	14519	1	2814	
14434	44802	Α	14520	1	3144	MGDFNTPLSTLDRSSRQKVNK
		1				DTQELNSTLHHADLIDIYRTLHP
]						KSTEYTFFSAPHHTYSKIDHVV
						GSKALLSKCKRTEIITNCLSDHS
						AIKPELRIKKLTQNRSTTWKLN
						NLLLNDYWVHNKMKAEIKMFF
						ETNENKDTTYQNLWDTFKAVS
						RGKFIALNAHKRKQKRCKIDTL
						ASQLKEVEKQEQTHSKASRRQ
						EITKIRAELKEIETQKTLQKINES
						RSWFLERINKIDRPLARLIKKKR
						EKNQIDVIKNDK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14436	44804	Α	14522	782	3024	SKTVPTGNSLEPTQNCWLQQW
						LPQDIEIEKQEQT/HSKASRRQEI
						TKIGAELEEIETQKTLQKINESR
						SWFFEKINKSDRPLARLIKKKR
						EKNQIDAIKNDKGDITTDPTEIQ
						TTIREYCKHLYANKLENLEEMN
						KFLDTYTLPRLNQEEVESLNRPI
						TGSEIEAIINSLPTKKSPGPDGFT
						AEFYQRYKEELHINRTKDKNH
						MIISIDAEKAFDKIQQRFMLKTL
						NKLVLEVLARAIRQEKEIKGVQ
1						LGKEEVKLSLFADDMIVYLENP
			:			IVSAQNLLNLISNFSKVSGYKIN
						VQKSQAFLYTNNRQTESQIMSE
						LPFTIASKRIKYLGIQLTRDVKD
						LFKENYKPLLNERKEDTNKWK
						NIPCSWVERINIVKMAILPKVIY
	:		:			RFNAIPIKLPMTFFTELEKTTLK
]		i		FIWNQKRARIAKSILSQKNKAG
						GITLPDFKLYYKATVTKTAWYS
						YQNRDIDQWNRTEPSEILPRIYN
	}					YLIFDKPEKNKQWGKDSLFNK
						WCWENWLAICRKLKLDPFLKP
						YTKIKSGWIKDLNVRPKTIKTL
			·			EENLGITIQDIGMGKDFMSKTP
				:		KAMATKAKIDKWDLIKLKSFC
						TAKETTIRVNRQPTEREKIFATY
		İ		1		SSDKGLISRIYKELKQIYKKRTN
						NLIKKWVKDMNRHFSKEDVYA
İ						AKKHMKKCSSSLAIREMQIKTT
]		MRYHLTPVRMAIIKKSGNNRITI
				ļ		VLLPGSLIVRSFHVTLNAILLHP
14437	44805	Α	14523	1	1701	
14438	44806	Α	14524	1	4434	
14439	44807	Α	14525	1	2913	MGGPAALCIAVVPGDLVVALC
						LEEGLGSGIATWACSPWRISQG
		1				THLTFRSSLAVAAIPEGLPIVVM
Ì						VTLVLGVLRMAKKRVIVKKLPI
						VETLGCCSVLCSDKTGTLTANE
						MTVTQLVTSDGLRAEVSGVGY
1						DGQGTVCLLPSKEVIKEFSNVS
						VGKLVEAGCVANNAVIRKNAV
						MGQPTEGALMALALKDQEDIY
		l				FMKGALEEVIRYCTMYNNGGIP
						LPLTPQQRGAGPGDTALPSAAS
						AFARMSAAERNDAFQG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
}	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14440	44808	A	14526	11	3009	MGDFNTPLSTLDRSMRQKVKK
' ' ' '	1	`	1.520	Î		DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIL
						GSKALLSKCKRTEIJTNYLSDGS
						AIKLELRIKNLTQNRSTTWKLN
						NLLLNDYWVPNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFIALNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
1						ITKIRAELKEIETQKTLQKINESR
						SWFFERINKIDRPLARLIKKKRE
				i		
14441	44809	 	14527	1	4800	KNQIDTIKNDK MGDFNTPLSTLDRSTRQKVNK
14441	44809	Α	14327	1	4800	-
						DTQELNSALHQADLIDIYRTLH PKSTEYTFFSAPHHTYSKIDHIL
						l .
						GSKALLSKCKRAEIITNYLSDHS
						AIKLELRIKNLTQSRSTTWKLN NLLLNDYWVNNEMKAEIKMFF
						ETKENKDTTYQNLWDAFKAVC
						RGKFIALNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQKTLQKINESR
						SWFFERINKIDRPLARLIKKKRE
11110	144010	<u></u>	1.4500	1	2001	KNQIDTIKNDK
14442	44810	B A	14528	569	2901 3443	RKH*TWKRTSSTSHHKNMPN*
14443	44611	A	14329	309	3443	LEKQEQTHSKASRRQEITKIRAE
						LKEIETQKTVQKINESRSWFFER SNKIYRPLSRLIKKKREKNQIDT
						IKNDKGDTTTDPTEIQTTIREYY
						KHLYANKLENLEEMDKFLDTY
				į		TLPRLNQEEIESLNRPITGTEIVT
	1					
						IINSLPTKKSPGPDGFTAEFYQR
						YKEELVPFLLKLFQSIEKEGILP
						NSFYEASIILIAKPGRDTTKKEN
14444	44012	-	14520	2501	0157	FRPISLMNIDAKILNKILAKRIQ
14444	44812	Α	14530	3591	8157	TGTSQKKTFMQPKKHMKKCST
						ITGHQSNANQNHNEIPSHTKLE
						WRSLKRSGNNRDVGNVVEAM
Ì					ĺ	YGDLPPPIMLIGHSMGGAIAVH
						TASSNLVPSLLGLCMIDVVEGT
				1		AMDALNSMQNFLRGRPKTFKS
}						LENAIEWRNIKDLSWADSIKNV
						AAASLACISGALPASAVTTSPD
				1		NKRMARNVLKYERLLAESPNH
						VVAEAVIQRPNIPHLQTRDTYE
[1		GLCQTLGSQPTLYQIPSLYCSYE
	<u> </u>	l				TNSNAYLLLQPIRKEVI

SEQ ID		Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	ŀ	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14445	44813	A	14531	172	328	AVKTEPEMKSQATKLSVVHEK
						KSQEGKPKEHTEPKSLPKQASD
						TGSNDAHNIKAVSR*SHRQPSC
s.						LWFMRKNPKKESRKNTQSQKA
						YPSRHQIQEVTMLTI
14446	44814	Α	14532	3	660	TSPDCWEEGRSVDWPRPGPALS
		ļ				LHNCKLDLGIEGGTLQPPTPSF*
		l				PTGTRAV/PTRRRSRQETQLNSE
		ŀ				RTVSPGSR**EKRTSRAW/RSPL
	1	İ				QGPGPGHAGQVAGGTGPRTGS
						TRGD/TLQKCGKIQSVPLNCELI
						PI*/PSILKPKVLPCSCPGIGASSS\
						CPSD*VFFCTS*PWPYLLSYPLS
						MNFYRYHDHILKLLVQYTLYL
						LLQFLQVLLKYLLSHPKQHS
14447	44815	Α	14533	1	1086	MVDNDVPWPLVVAQEKGQPD
						LCLSSKGYRPLALVQLDFHQVS
						VTSKIFILSGALRSLTGAATAD
		ŀ			·	WTCRVLVIYPHDIMADSECNSL
		ŀ				LVPSTTRKIMKHMLASEKIKTQ
						NFNYNVDECVSLLYHHKANKL
						EAAEARQECELLRVASVSRHC
 -		İ				W/RDPAHPHSCWPRC*APHCLG
ŀ						PARAPNARPAKPATTWNLRSSR
		İ				APCAAPVPTQSWAGAPSGTAG
						AEPS*ECRPPENQARTAGSGVH
				ŀ		AEACAPATCPACPGPGPCSGLP
						SSGRALFLL/CATLAIPCV*V*LC
						FLATPPPCR*HVTAPSAGKCIW*
						PDA*VTK/PRDLCGPVQVARDN
						GPRQVAQKKESSSYVSWFGTFI
						SRVGDRSPPPRRAATL
14448	44816	Α	14534	23	506	PDIDNIFSEFGAKSKPYLTVDQ
						MMDFINLKQRDPRLNEILYPPL
						KQEQVQVLIEKYEPNNSLARKG
						QISVDGFMRYLSGEENGVVSPE
						KLDLNEDMSHPLSHYFINSSHN
				1		TYLTGMNFLVLCYFSQSHQISP
				1		TSTFCCSEAEV/WFSSISCEVPLR
						TLGFHH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ŀ		sequence		
14449	44817	Α	14535	341	1248	KNAFTSFLTSFPPGQISVDGFMR
						YLSGEENGVVSPEKLDLNEDM
						SQPLCFMGIWGLFCWCFIAGQL
						AGNSSVEMYRQVLLSGCRCVE
						LDCWKGRTAEEEPVKILAY*IY
						MCSHF*EVIEAIAECAFKTLSLP
						TFLLLFLFSLSPKQQAKMAEYC
						RLIGSSIFKEYAFHFY*LESGVPL
				1		PSPMDLMYKILVKNKKKSHKS
						SEGSGKKKLSEQASNTYC*NGD
						MMICY*NLTGEADTESD\DADS
						GNV/CLFYSYFQGTAGSEAMAT
						EEMSNLVNFLTFPFFVLLLLNSE
						RNKSFEMSSFVETKGL
14450	44818	Α	14536	574	647	C*PGNM/HISGQHLVAGEKEAE
				-	:	SEDEEEEDMKLLGTPEGGSKFT
			•			QKKVKLAADEDDD/DDDADDD
				•		DDDDDDDDDDDEETEEKAPV
14451	44819	Α	14537	3	630	PGATHASAPPGVLRLKCGSGPV
					1	HIIGQHLVAVEEDAQSQDEEEE
				-		DVKLLSISGKRSAPGGGSKVPQ
				1		KKVKLAADEDDDDDDDEEDDD
		1				EDDDDDDFDDEEAEQQAPVKK
						SI\RDTPAKNAQKSNQNGKDSK
						PSSTPRSKGQESFKKQEKTPKTP
		}				KGPSSVEDIKAKMQASI/EKKG
				İ		GSLPKVEAKFINYVKN\CSRMT
						DQEAI\QDLWRWRRSL
14452	44820	A	14538	1	1971	DUGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG
14453	44821	Α	14539	316	1169	RVGGQSHGTQRISLCRHSVCSP
						LARQSPSNMK*VRTEEIQMAVS
				1		CYLKRWQYVDSDG\PLKQGLR
						LPQTAEELAANLTVHRESHCAD
						IVSAVP\CQAEPQQYEV\QFGRL
						RNFLIDSDSPHSHEVMPLLYPLF
						VCLHLSLVQNSPKSTVESFYSR
			į.			FHGMFLQKAS\QKGVIEQLQTT
]					QTIQDILSNFKLRAFLDNNYVV
:						HLQEDSYNYLIHYLQSDSYTAL
						CKMFILLIHLDVQPAKTTD*QL
						YASGSSSRSENNYLEAPDMPSPI
						LQNDAALEVLQERIKQVKDGC
						PSLTTVCFYNT*QLLNTAEISPD
						SKLLAAGFDNSCIKLWSLQSRK
1112	14000	ļ. —	14540	ļ <u>.</u>	207	LKSESCQVDMSRIHLACDILEE
14454	44822	A	14540	1	297	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide inscrtion)
14455	44823	A	14541	2	1089	APLLGECRSGRGLASSRSRRRT TMTQSVVVQGKLRRAGRDLVA AESSGHVSCLMSYSLSFLSSGRS RLALREHAAVNQVPARRREPPR LSARSSSPYSFHLQSE*L*KKIH* LYFIFLSQAVLIDMEEGVVNEIL L*IRTNQIKK*MQGIFKIHILIFR AVGHKVFGSLYQDQILEKFQIL HNYILNHLC*FLKTL*SSGTGSG LGTFLLKVLEDEFPEVYRFVTSI YPSGEDDVITSPFLVEYHLHPLF LIASLPMFFPQKSLFDIISKIDLM \VNSGKV/GGTTVKPKSL\VTSSS GALKIELHLKCIILKLKIFSF*C* LFSSARFEGSLNMDLNEISMNL VPFPPSS*EIVSLKYVLYFN
14456	44824	A	14542	129	461	TCISGWHLFRASVTCPGIQTSPH NWHISDNSSLPNNIFIVCWNLL GMTRNYIIFIEQPLKMNLWKIA TSKIRGKAFSDGIS\GNPQCNTR FHVVEKRTGQLLPGRYYSKPF
14457	44825	Α	14543	1	522	
14458	44826	A	14544		927	MEDLLTEARSPYWGRARAKIQ DSCWSVTSDPNAGSVDILAPLL LPSPSSYLPNFTCTVSVVTSHLH EAFLDLGFRGTGGSNCKTSRQV QLNNQKRKKKKSKSLPFHMSL QIQNYKTQEEPHAKPSSQQHQH SEHESIQTKLVFRSSMTYMHTL KLIREKGSIMLSSTQGECPNSTC RWPAEEHPLGSTHREDQEHVIR VPPEKVDLGETIHGVQVICSIAS TEKGKPSYYHSFGMTRNYIIFIE QPLKMNLWKIATSKIRGKAFSD GISWEPQCNTRFHVVEKRTGQT T*INPDKISIQKQHDIHAYIEAD KRKRKHNAVIHTG*M/YQI/GTC RWPAEEHPLGSTHREDQEHVIR VPPEKVDLGETIHGVQVICSIAS TEKGKPSYYHSFGMTRNYIIFIE QPLKMNLWKIATSKIRGKAFSD GISWEPQCNTRFHVVEKRTGQNI LNVKTMRMKTFMMIYYHLMN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14459	44827	A	14545	1	882	MAEILTQVPDDMLKKKTPRVK
						SVWRSQRGSCTPLIGTTELTLD
ŀ						TSHMSIRNMDRSHINVHTVRKP
						SADLPYFRTHEWAHTGGKPYD
						CEECGKSFISRSSIRRHRIMHSG
						DGPYKCNFCGKALMCLSLYLIH
						KRTHTGEKPYECKQCGKAFSH
Ì						SGSLRIHERTHTGEKPYECSECG
ĺ						KAFHSSTCLHAHKITHTGEKPY
						ECKQCGKAFVSFNSVRYHERT
		İ	†			HTGEKPYECKQCGKAFRSASH
		ŀ				LRTHGRTHTGEKPYECKQCGK
						AF*PRFQMTC*RRKLPE*NQCG
						EVSVGH\HSLNRHHRADTGHKP
						YEYQEYGQKPYKCTYCKKAFS\
						YLPYFRTHEWAHTGGKPYDCE
						ECGKSFISRSSIRRHRIMHSGDG
						PYKCNFCGKALMCLSLYLIHKR
1						THTGEKPYECKQCGKAFSHSGS
						LRIHERTHTGEKPYECSECGKA
						FHSSTCLHAHKITHTGEKPYEC
						KQCGKAFVSFNSVRYHERTHT
						GEKPYECKQCGKAFRSASHLRT
	;					HGRTHTGEKPYECKQCGKAFG
						CASSVKIHERTHTGEKPCSSNTS
14460	44828	Α	14546	3	395	SAEVGAAETTLTELRRTVQSLEI
		ļ.				DLDSMRNLKASLENSL\GILLHL
						ESELAQTRAEGQRQAQEYEAL
						LNIKVKLEAEIATYRRLLEDGE
						DFNLGDALDSSNSMQTIQKTTT
						RRIVDGKVVSETNDTKVLRH
14461	44829	Α	14547	2	1044	AKTSSQMPSPQEGMCGKACTP
						ALSQADSLCPLLRLASEVEGYS
						LPACAEPYVQSECLSHLSVWSL
						QHALSSLASMSFTTCSAFTNYW
						SPGSVQVPSYGTQPVSHAASVY
						AGLGGSGSRISVSHFTNG\GGLA
						GMGGIQNEKETMQSLRDRLAS
						YLDRVRGLETENWKLESKIQEH
						LNVTRL*LETEIEALKKELLFM
						KKNNEEEAKGLQAQIASSGLTV
						EVDAPKS/QDLAKNMADSRAQ
						YDKLARKNREELDKYWSQQIE
						ESTRVVTTQSAKVGAAEMTLT
						ELRHRVQSLEIDLDSTRNLKAS
						LENSLREVEARYALQMEPSLLF
		<u></u>				SKIIFGRVWEISYLHVFEGKQV

	of peptide	١	ı			
· I		hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
. 1	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14462	44830	A	14548		1212	<u>GPGTGIAGGLAGMGGIQNEKET</u>
14402	44030	\^	14340	1	1212	MQSLNDRLASYLDRVRSLETE
						NRRLESKIREHLEKKGPQVRD
						WSHYFKIIEDLRAQIFANTVDN
						ARIVLQIDNARLAADDFRVKYE
		ŀ				TELAMROSVENDIHGLRKVIDD
						TNITRLQLETEIEALK*ELLFMK
						KTPEEEVKGLQAQIASSG\LTVE
						V\DAPKSQDLAKIMADIR\AQY
						DELARK\NREELDKYWSQQIEE
		ŀ				STTVVTTQSAEVGAAET\TLTEL
						RRTVQVFGDSTWTSMRNLKA\
		ļ				NLENSL\GEVEARYAL\QMEQL\
						NGILLHLESQLGQTPRTEAQRQ
						AQE\YEALL\NIKVKL\EAEICHL
						TRPPSWKIGEDFNL\GDS\LDER
						NSMQTIQKTTTRRISWIGQSGV
						LRPIDTKVLEALSQQEASGTLL
14462	44001	_	14540		2292	GKQEANKKFQSFK
	44831	A	14549	1	2382	
	44832 44833	A C	14550	2	173 732	
	44833	A	14551 14552	455	682	AWKRAVALGKSRPMGAVAFK
14400	44834	A	14332	433	082	RRMQPVAGPRAMQEKPFPSSSS
						SLSIPPAF*YRLALSLAEPSRRW
			:			RARTGALMQL
14467	44835	Α	14553	158	368	IN HET GITEMQE
	44836	A	14554	87	251	
	44837	Α	14555	1	1026	
	44838	A	14556	201	578	
	44839	A	14557	336	767	AFVRAGSRFGGTPGARPDPAPA
						TRAPNPVGQSRHLRAASSRGRL
						FLRRTRDRKFPRRPRLSRRLLG
						AI/GCRWNQPRAAEKSFSAGAV
						LDFSPGLSCLPAVQGSGPAARH
						A*ASHPLHGLLCCPSIPDEHHPL
						LHGAQSHRPPKG
14472	44840	Α	14558	6	535	ARHRVLIGVFTIPELDILSTSPPD
						SGAQLASPS\DPAPGLQVELPAS
						PARCARIPOPLGGRWDWAQTQ
						EPSWLHLVEPAPGLQVELPASP
						TPCACIPQPLGDRWDWAPWSR
						GWCSLGRLGPHRSPWSGWEAE
						AWRAAGPEPCPTGRQLRLGEK
						SSAAPVGWHCWGTQYTLRSH
						WPGC
,		<u> </u>			1600	
14473	44841	lA –	14559	375	638	

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
14475	44843	Α	14561	54	380	SCHHYCRIHRRCARFLAGP*LPS RGARLGTCSPPCVSLPPAPWAP VRPEPPRRAPPPAPQRPVPSTTQ GLRSAGARRGTGRQLHLQLRG SAASFLKSVRPRTHQFRTH
14476	44844	Α	14562	531	773	
14477	44845	В	14563	92	1774	
14478	44846	A	14564	1	390	LHDLQLSKPLSDDTALTDGG/Q PLPARWWAQPGRLQRCARFLA GP*LPSR/WGRARDLQPAMPEP PTPSVGSCAAQASPTSAAPCSM APSPIDHPRAWPVLCSFCMLLS LFVASISSCSQHQPMQVGTMVC
14479	44847	Α	14565	165	545	SLHSKLCWWLQQLWVTDTCPR CSVQDCRQRWASAGNQS*RLL VDPAPGLQVELPANPVPCARIP QPLGGRWD*APWSRGWCSSGR LGLHRSPWSGWEAQAWRAAG PEPRPAGRQLGAMSCKVETGT
14480	44848	A	14566	I	309	SACLAGRPIRGEGRPRDLQPAM PEPPTPSMGSCAARASPTSATPC STAPSPIHHSRAEECGHTALG/H GRQLHLQPRNLPQTINWRTTYS SFPATLNRLDGGG
14481	44849	В	14567	1	843	
14482	44850	Α	14568	143	535	
14483	44851	Α	14569	375	571	
14484	44852	В	14570	289	972	
14485	44853	Α	14571	27	363	
14486	44854	Α	14572	2	490	
14487	44855	A	14573	574	961	RLTLPDRLGSPPDTHQAQQITW ALLPQGFADSPHYFSQAQISSSS ITYLGIILHENTRALPADHV*LIS QTPISSTKQQLLSFLGMVRYFC LWIPSFTILTKPLY*FTKANLAD PTDPKSFPHSSFRSL
14488	44856	В	14574	182	1513	
14489	44857	A	14575	894	1305	QTQEPSWLHLVDPALGLQVELP ASPVPCARIPQPLGG*WDWAP WSRGWCSSGRLGPHRSPWSGW EAQAWRAAGPEPCPAGRQLRP GQHPL/RHMNKSQALSLLIAQP LGINLLSWWPPKYNRPVPLIGL SGLLIPATF
14490	44858	Α	14576	531	801	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14491	44859	Α	14577	441	783	RAPPPAPWRPVPSTTQGLRSAG
	!		:			EQRGTGRQLHLSAAPVRDPRG
						EAIWAPESGRSAASILKPARPPA
						HREERTTPDAPP*EL*HSPRRSA
		1				ASLREDPWLHS*SQ*DQEPTNS
				-		GHTVSQDQSYYS*LEVSETKNP
						PIPDTL
14492	44860	Α	14578	344	1632	FLAGPYLPSHGAGLGTCSLPCL
			•		•	SLPPTPWAPVRPEP/PPMSTTPCS
						TAPSPIDHPRAEECERTAQDWQ
						AAPPAAPVWDPLGEASWAPES
				ŀ		GTNISITPGNATFVTRVQGQAW
						FASCITGHDMSSLKTSSVVVLR
						RQSEALLPANLTCNWQDSSAL
:					:	ATLESALSQVRHKRFTFTLMVF
					1	TGSAIDIMATVSIAVVSITESVQ
						TTAFVDNLAKNICDELLLQDM
						DEAGNQHSQQNNTRTENQTPH
						VLTHKWELNNENIRIQGGEHYI
						LGPVGGRDEKTLLLRTYCLKSQ
}						EGADTKCRNKHNKETQKQSKS
						LVYSSTLNPEKLLVCKRLSLEL
						TRPENTLRCTNIGKLKAVAGG
						MSAASGIYLETDTQPYPLKQEL
ł						KEIKECVGKNSVVVVFLVRKK
		1				LSVSQYAFARCLDAKHSTSENV
						EIAGKLRIQSFEVGKQL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14493	44861	A	14579	3	2233	NWLPSVSRCQETHKIAANQSLT
		l				QRQMPLTSLYCDVVILPVSVGL
						ESLQSPYYRTSPCDPVSQFCLIK
						ISASSLPVLSLWKTLKNAQSYQ
						RDYRHDNKAHYFQKLVTMQE
						NLYVLLCDEELDAVSLEAVHQ
İ						QKMKNRQAHSSSGCCQKVAN
1						APRNEESSGQFRVIGNEKFEKG
1						PCGVRQQIFEKVLLPNRTCEFV
						TPRFNRNFEDLIKDLKAAETLID
						FLVYEPKNQRRLVNRFRLPCKL
						LHNLGHPLKTDAAQSPRKPPGP
		i				SWMPSFGSKPPSHPPLVSPHLNP
		l				QVWDTSSPSLATEHASLTISLKP
						NHPYPAQSQYSIPQHALKGSKP
						VITRLLEHGLLKPINSPYNSPILP
						VQKLDKSYRLVQHLCLIYQIVL
ľ						PIHPMVPNPYTLLSSIPFTTHYS
						VLDLKHAFFTIPLYPSSQPLFAF
İ						TWTDPDTHQAQQITLAVLPQCF
						IDSPHYFSQAQISSSSVTYLGIIL
						MKTHIGLGAVEQGVVLVGEAR
						AAQEPMEWVGGSGM/CGLQVP
1						SPAPWEGS*GPARNRAQ\PVTIV
						PVLDFNPAFHIIPDTTPDHHDCI
						SLIHLTFTPFPHISFFPVPHLEHT
	1					WFIDGSSTRPNCHSPAKAGYAI
						VSSTSIIEATALPRSTTSQQAKLI AYTDSKCAFHILHHHAVIWEER
						NFLTMQGASIINAFLIKTLLKDI
						LLPKEAGVIHCKGHQKASDPIT
		<u> </u>				QDNAYADKVAKKQLAFQLLSL
14494	44862	Α	14580	1	642	SGTVPPTFVRTVIQAQGARTVI
						QQSLIGSLYIFRHCVRCQGIESV
						FVIIIWSRVGYDEKGFCLFTSVL
			1			GVWGHVRTKPCLLITWRTLKS
						RSGLE*PAGFSSSPGTSPQIIRKQ
						KRGVPQHPVCDLPGRFRPRAGT
			1			PLAAQIQVPGLPPCGHQE*NRR
]			WAGSSALCPPYLCARAGTGAV
	1	1	1			CRACEPAGV/AGWAWAWWAP
ļ		<u> </u>				HSEHAASPAGPRQWGT
14495	44863	В	14581	393	549	
14496	44864	В	14582	1	591	
14497	44865	C	14583	226	828	
14498	44866	Α	14584	1	1362	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence	l	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14499	44867	A	14585	1	585	MANVHQTMPMRLRKLPDSFKP PEPKSNSQRVSTEAGTAGALTP QHVRAHSSPASLQLGAVSPGTL TPTGVVSGPAATPTAQHFQQSS FEIPDDVPLPAAREMAKTSSGQ RYFLNHIDQKTTQQDP\RKTML SQMNVIANPPVQQNMMNS\PQS PQGGV/MGGSNSNQQRQMRPQ QLQMEERLQLKQQ*LLQQELA LR
14500	44868	A	14586	345	1053	RWRLQNLSFCSSCSGSGSEGAT CFARGRQQSNLQSKIKFPSNTD SRSTGYNSFDRQPNICFYKRQE TPHTGPHPVPEGRPYPMPG*CG A*SPGPFASRN*SEKN*SYKCSL HTAFNVIFSSVI**RYCTRQW\HI VKCLDSFCDPFLISDELRRASTD AGTAGALTPQHVRAHSSPASLQ LGAVSPGTLTPTGVVSGPAATP TAQHLRQSSFEIPDDVPLPTSW GEAKTSSGSKIPF
14501	44869	В	14587	607	1185	

1818

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14502	44870	A	14588	1	2142	MIILIDAEKAFDKIQQPFMLKTL
						NKLGIDGTYLKITRAIYDKPTA
						NIILNGQKLEAFPLKTGTRQGCP
						LSPLLFNIVLEVLAQAIRQEKEI
						KGIQLGKEEVKLSLFADDMILY
[LENPIVSAQKLLKLISNVSKVSG
						YKINVQKSQAFLYTNNRQTESQ
Ì						IMSEFPFTIATKRIKYLGIQLTRD
						VKDLFKKYKPLLNKIKEDTNK
						WKNIACSWIGRINIMKMAFPR
						WELNNENTWTQEGEHHTLGPV
						VGWGKRGGIALVDIPNVNDKL
						MVLEVLARAIRQKKEIKGIQLG
						KEEVKLSLFADDMIVYLENSIV
						SAQNLKLISNFSKVSGYKINVQ
						KSQAFLYTNNRQTESQIMSEFP
						FTIATKRIKYLGIQLTRDVKDLF
						KENYKPLLKEIREDTNKWKNIP
						CSRIGRINIMKMAILPKVIYRFN
						DIPIKLPMTFFTELEKTTLKFIW
						NQKRACIAKTILSKKNIAGGITL
						PDFKLYYKATVTKTAWYWYQ
						NRDIDQWNRTEASEVTSHIYNH
-						LIFYKPDKNKKWGNDSLFNKW
						CWENWLAICRKLKLDPFLTPYT
						KIHSRWIKDLNVRPKTIKTLEEN
						LGNTIQDIGMGKDFMTKTPKA
ļ			1			MATKAKVDKWDVIKLKSFCTA
				·		KETTIRVSRQPTEWEKIFAIYPS
ł						DKGLISRIYKELKQIYRKK\TNN
				,		PIKKWAKNMNRHFSKEDIYAA
						NRQMKKCSSSLVIREMQIKTTM
14503	44871	В	14589	1	1647	
14504	44872	В	14590	1	2025	
14505	44873	В	14591	4	2121	
14506	44874	Α	14592	2	1879	
14507	44875	Α	14593	1	2130	
14508	44876	Α	14594	1	3018	
14509	44877	В	14595	1	3171	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14510	44878	A	14596		2700	MEKKQSRKTGNSKNQSASPPP
1 15 10	11070	1	1.570			KECSSLQATEHSWMENDFDEL
						REQGFRRSNYSELKEEVRTHGK
						EVKNLEKRLDEWLTRITNAEKS
						LKDLMELKTMARELLDECINLS
						SQFDQLEERVSVMENQMNEMK
	ĺ					PNLRLIGVPESDGENGTKLENT
						LQDIIQENFPNLARQANIQIQEIQ
						RMSQRYSSRRAIPRHIIVRFTKV
		1				EMKEKMLRAAREKGRVTLKG
						KPIRLKADLSVETLQARREWGP
:						IFNILKEKNFQPRTSYPGKLSFIS
		l				EGEMKSFTDKQKLRYFVTTRP
						ALKELLKEALNMERNNRYOPL
						EKQIQTTIREYYKHLYANELEN
				ļ		LEEMDKFFDTYTLPRLNQEEVE
						SLNRPITGFEIEAIINSLP/TKKSP
						EPGGFTAEFYQ\RIEKEGILPNSF
Ì						YEASIILMLKPGRDTTKKENFRP
						ISLMNIDAKILNKILANRIQQHI
						KKLIHQDQVGFIPGMQGWFNIR
						KSINIIQHINRTNDKNHKIISIDA
						EKAFYKIQQPFMLKTLNKLGID
						GSYLKIIRAIYDKPTANIILNGQ
						KLEAFPLKTGTRQGCPLSPLLF
						NLVLEVLARAIRQEKEIKGIQLG
						KEEVKLSLFADDMIVYLENPIA
						SAQNLLKLKGNFSKVSGYKINV
						QKSQAFLYTKNRQTESQIMSEL
						PFTIASKRIKYLGIQLTRDMKDL
						FKENYKPLLNEIKEDTNKWKNI
						PCSWEGRISILQMAILPEVIYRF
14511	44879	A	14597	1	2310	
14512	44880	Α	14598	1	1305	
14513	44881	A	14599	1	2148	
14514	44882	В	14600	1	777	
14515	44883	A	14601	2	2971	
14516	44884	A	14602	1	3810	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14517	44885	A	14603	1	1773	MNIGAKIINKILANQIQQHIRKLI
						HHDQVGFIPGMQGWFNICKSIN
						VIQHINRTKDKNHMIISIDAEKA
						CDKIQQRFMLKTLNKLGIDGKY
						LKIIRAIYDKPTANIILNGQKLE
						AFPLKTGTRQGCPLSPLLFNIVL
			1			EVLARAIRQEKEIKDIQLGKEEV
						KLSLCADKMIVYLENPIVSAQN
						LLKLISNFSKVSGYKINVQKSQ
						AFLYTNNRQTESQIMSELPFTIA
						SKRIKYLQIHQTRDVKDLFKEN
						YKPLLKEIKEDTNK\NIPCSCVG
				E		RINIVKMAILP/KELEKTTLKFIW
						NQKRARITKSILSQKNKAGSIM
ı					}	LPDFNLYYKATVSKTAWYWY
		ĺ				QNRDIDQWNRTEPSEITPHIYN
			ļ			YLIFDKPEKNKQWGKDSLFNK
						WCWENWLAICRKLKLDPLLTL
						YTKINSRWIKDLNVSPKTIKTLE
1		1			İ	ENLDITIQDIGMGKDFMSKTAK
1		1				AMAAKAMSTKAKFGKWDLIK
		i				LKSFCTAKETTIRVNRISSYASS
1						STSLQAFSRKILNCSKRFFKIAQ
1						MCKLVELRPQDMHTSRGSGNC
					1	SEARTLLCDFSAGRTSVNHGRI
						WRQIREAGAADPEIGLEVCNKN
						PTSFTLFYFLSYTTG
14518	44886	В	14604	1	2070	
14519	44887	В	14605	1	3190	
14520	44888	Α	14606	1	3370	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
			<u> </u>	sequence		
14521	44889	Ā	14607	569	3030	RKH*TWKRTSSTSHHKNMPN*
						LEKQEQTH/SKPSRRREITKIRA
						ELKEIETPKTAQKINESRSWFSE
						RINKIDRPLARLRKKKREKNQI
						DTIKNDEGDITTHPTEIHTIIREY
						YKHLYANKLENLEEMDKFLDT
						YTLPRPNQEEVESLNGPIAGSEI
						QAIINSLPTKKSPGPDGFYQRYK
						EELVPFRLKLFQSIEKEGILPNSF
						YEASIILIPKPGRDTTKKENFRPI
						SLLNINAKILNKILANRIQQHIK
						KLMHHDQVGFIPGMQGWFNIR
:						KSINVIQHINRTKDKNHMIISID
						AEKAFDKIQQPFMLKTLNKLGI
						DGTYHKIIRANYDKPTANIILNG
						QKLEAFPLKTGTRQGCPLSPLL
						FNIVLEVLARAIRQEKETEGIQL
1						GKEEVKLSLFADDMIVYLENPI
						VSAQNLLKLISNFSKVSGYKIN
						VQKSQVFLYTDNRQTESQIISEL
		İ				P\FTIASKRIKYLGIQLTRYVKDL
						FKER/YNEIKEDTNKWKNIPCS
						WVGRINIVKMAILPKVIYRFNAI
						SIQLPMTFFTELEKTTLKFIWNQ
						KRAHIAKSILSKKNKAGGIMLP
						DFKLYYKATVTKTAWHWYQN
						RDIDQRNRTEPSEIMPHVYNHLI
						FGKPDKNKQWGNDSLFNKWC
						WENWLAICKKLKLDPFLTPYT
						KINSRWIKDLNVRPKTIKTLEEN
						LGNNIQDIGMGKDFTSKTPKA
1.000	1,1000	<u> </u>	14600		0740	MATKDKIDKWDLMKLKSFCTA
14522	44890	A	14608	1	2742	
14523	44891	A	14609	1	3828	
14524	44892	A	14610	1	2667	
14525	44893	A	14611	1	2325	
14526	44894	Α	14612	1	3057	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14527	44895	A	14613	603	2973	DNTARGTIRQQHLLFTNIRCSA
1,132,	11055	l' .				ASAADTQANRVWSGPPANSNR
	,					PAA\RVLTVRRKTNKQKGHPH
						QNPICTSPSSKTKETQTTIREYY
						KHLYANRLENLEEMDKFLDTY
						TLPRLNQEEVESLNRRITESDIE
						AIINSLPTKKSPGPDGFTAEFYQ
						RYKEELVPFLLKLFQSIEKEGIL
						PNSFYEASIILIPKLGRDTTKKE
						NFRPISLMNIDAKILNKILANRI
		·				QQHIKKHIHGQVGFIPGMQGW
						FNIRKSINVIQHINRTNDKNHMI
						VSIDAEKAFDKIQQPFMLKTLN
		•				KLGIDGPYLKIIRAIYDKLTANII
						LNGQKLEAFPLKTGTRQGCLLS
						PLLFNIVLEVLARAIRQEKEIKGI
						QLGKEEVKLCLFADDMIVYLE
1						NPIVSAQNLLKLISNFSKVSGYK
						INVQKSQAFLYTNNRQTESQIM
						SELPFTIASKRIKYLGIQLTGDV
						KDFFKENYKPLLNEIKEDTDKW
						KNIPCSRVGRINIMKMAILPKVI
						YRFNAIPIKLPMTFFTELEKTTL
						KFIWNQKRARIAKSILSQKNKA
						GGITLPDFKLYYKATVTKTACH
						RVGRAQQHSISKWKWYIHDWS QVGPEGTNDSARYPDTTQKWT
						AAALQPLSRTSLKDSHEGKSSQ
						WAELRAVHLVLRFAWKEKWP
						DVQLYTDSWAVASGLAGWSG
						TWKKHDWKIGDKEIWGRATPV
						IAQWAHEQRGHGGRDGDYAW
14528	44896	В	14614	1	3105	
14529	44897	A	14615	1	3654	
14329	17707/	Λ	14013	1	3034	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14530	44898	A	14616	735	2839	RSFPRSRPSPFLLLSRYLRIHMV
						FSVLPFGLQNPKYLLSGSLQEK
		l				FRTPGINSHKTLDPNRVIIKVRR
						RLEKEKALRAYVGKSEVRNMH
						LRERCKL*KKREKNQIDTIKND
						KGDITTDATEIQTTIREYYKHLY
:						ANKLENLEEMDKFLDTYILPRL
		1				NQEEVESLNRPITGSEIEAIINSL
		ļ				PTKRSPGPDGFTAEFYQTYKEE
						LVPFPLKLFQSTEKEGILPNSFY
						EASIILIPKPGRDTTKKENFRPIS
		1				LMNIDAKILNKILANRIQQHIKK
						LIHYDQVGFIPGMQGWFNKRK
						SINVIQHINRTNDKNHMIISIDAE
						KAFDKIQQPFMLKTLNKLGIDG
						TYLKVIRAIYDKPTANIILNGQK
						LEAFPLKTDTRQGCPLSPLLFNI
						VLEVLARAVRQEKEIEGIQLGK
						EEVKLSLFADDMIVYLENPIVS
		1	}			AQNLLKLISNFSKVSGYKINIQK
1		1				SQAFLYTNNRQTESQIMSELPFT
						TASKKIKYLGIQLTRDVKDLFK
		l	•			ENYKPLLNEIKEDTNKWKNIPC
		l				SWVGRINIMKMAILPKVICRFN
		ł				AIPIKLPMPFFTELETTTLKFIWN
		l				QKRARIAKSILSQKNKAGGITLP
		İ				DFKLYYKATVTKTAWYWYQN
						RDIDQWNRTEPSEITPHIYNYLI
		ļ				FDKPEKMSIIDTGGWYEAPVSS
						FFKEGLCRTGTENQVNRQPPAA
						SPFKVYVSCTGSLHPRPHYFLE
14531	44899	Α	14617	1	2250	
14532	44900	Α	14618	1	2870	MKAEIKVFFETNENKDTTYQNL
		•			,	WDTFKAVCRGKFIALNAHKRK
		ľ				QERSKIDTLTSQLKELEKQEQT
						HSKASRRQEITKIRAELKEIQTQ
		}				KTLQKINESRSWFFERINKIDRS
						LARLIKKKREKNQIDTIKNDKG
						DITTDPTEIQTTIREYYKHLYAN
						KLENLEEMDKFLDTYTLPRLNQ
1		1				EEVESLNRPITGAEIVAIINSLPT
		1				KKSPGPDGFTAEFYQSWAETQP
}						KKENFRPISLMNIDAKILNKILA
						KRIQQHIK

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14533	44901	A	14619	248	2623	RQWAGVVGRCSHLASWVSSNT
1						SETGAIRSSTEVDAPDDSMLLST
						CDIDLTAARRAWLGCLPTKKSP
						GPDGFTAEFYQRCKEELVPFLL
		İ				KLFQSI/EKEGILPNLFDEASIILI
						PKRGRDTTKKENFRPISLMNID
						AKILNKILANRNQQHIKKLIHH
						DQVGFIPGMQGWFNICKSINVI
						QHINRTKDKNHMIISIDAEKAF
						DKIQQPFMLKTLNKLGIDGTYL
						KIIRAIYDKPTANIILNGQKLEAF
				1		PLKTGTRQGCPLSPLLFNIVLEV
				ļ		LARAIRQEKEIKGIQLGKEEVKL
			ļ	i		SLFADDMIVYLENPIVSAQNLL
				i		KLISNFSKVSAYKINVQKSQAF
						LYTNNRQTESQIMSVFPFTIASK
						RIKYLGIQLARNAKDLFKENYK
						PLLNEIKEDTKKWKNIPCSWVG
						RINIVKMAILPKVIYRFNAIPIKL
						PMTFFTELEKTTLKFIWNQKRA
				:		RIAKSILSQKNKAGGITLPDFKL
						HYKITVTKTAWYWYQNRDIDQ
				1		WNRTEPSEITPHIYNYLIFDKPE
						KNKQWGKDSLFNKWCWENWL
1						AICRKLKLDPFLTPYTKIHPRWI
						KDLNVRPKTIKTLEENLGNTIQ
						DIGMGKDFMSKTPKAMAAKA
				Ī		KIDKWDLIQLKSFCTAKETTIRV
						NRQPTKWEKIFATYSSDKGLIS
		ŀ]		RIYKELKQIYKKK\TNNPIKKW
		İ				AKDMNRHFSKEDIYAANKHM
						KKCSPSLAIREMQIKTTMRYHL
14534	44902	Α	14620	1	3253	MGDFNTPLSTLDRSTRQKVNK
					Ì	DTQELNSALHQGDLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
İ						GSKALLSKWKRTEIITNYLSDH
						SAIKLELRIKNLTQSRSTTWKLN
						NLLLNDYWVHNEMKAEIKMFF
			1			ETNENKDTTYQNLWDAFKAVC
	1.					RGKFIALNAHKRKQERSKIDTL
					1	TSQLKELEKQEQTHSKASRRQE
			1			ITKIRAELKEIETQKTLQKINESR
						SWFFERINKIDRPLARLIKKKRE
						KNQIDTIKNDK
14535	44903	В	14621	1	2758	
14536	44904	A	14622	1	5178	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
			<u> </u>			
14537	44905	Α	14623	1	3382	MGDFNTPLSTLDRSTRQKVNK
						DTQEFNSALYQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
·						GSKALLSKCERTEIITNYLSDHS
						AMKLELRIKNLTQNCSTTWKL
			İ			NNLLLNDYWVHNEMKAEIKM
						FFETNENKDTTNQNLWDAFKA
						VCRGKFIALNAHKRKQERSKID
						TLTSQLKELEKQEQTHSKASRR
						QEITKIRAELKEIETQKTLQKIN
						ESRSWFFERINKIDRPLARLIKK
						KREKNQIDTIKNDK
14538	44906	Α	14624	965	4091	TWKGTTSTSRCKIMPKYRSTRQ
						KVNKDTQELNSALHQADLIDIY
						RTLHTKSTEY/TFFSAPHHTYSK
						IDHIVGSKALLSKCKRTEIITNY
						LSDHSAIKLELRIKNLNQSRSTT
						WKLNNLLLNDYWVHNEMKAE
						IKMFFETNENKDTTYQNLWDA
					}	FKAVCRGKFIALNAHKRKQERS
		1				KIDTLTSQLKELEKQEQTHSKA
						SRRQEITKIRAELKEIETQKTLQ
						KINESRSWFFERINKIDRPLARLI
						KKKREENQID
14539	44907	Α	14625	3	3229	
14540	44908	В	14626	1	2791	
14541	44909	Α	14627	1	3158	MVKGSIQQEELTILNIYAPNTG
						APRFIKQLLSDLQRDLDSHTLIM
						GDFNTPLSTLDRSTRQKVNKDT
		ļ				QELNSALHQADLIDIYRTLHPK
				j		STEYTLFSAPHHTYSKIDHILGS
						KALLSKCKRTEIITNYLSDHSAI
						KLELRIKNLTQSRSTTWKLNNL
						LLNDYWRKQERSKTDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
						AELKEIETQKTLQKINESRSWFF
						ERINKIDRPLARLIKKKEKNQID
						TIKNDKGD
14542	44910	В	14628	1	7849	LANGE TO PORT A PORT OF THE COLUMN ASSET
14543	44911	Α	14629	Į I	3985	MENDFDELREEGFRRSNYSEL
						WEDIQTKGKEVENFEKNLEECI
						TRITNTEKCLKELMELKTKARE
		1				LHEECRSLRSRCDQLEERVSAM
						EDEMNEMKGEGKFREKRIKRN
		1				EQSLQEIWDYVKRPNLHLIGVP
		1				ESDGENGTKLENTLQDIIQENFP
				}		NLARQANVQIQEIQRMPQRYSS
		1	1			RRATPRHIIVRFTKVEMKEKILK
		1				AAREKDRSTRQKVNKDTQELN
						SALHQADLIDIYRTLHPKSTEYT
			<u> </u>	<u></u>		FFSAPHHTYSKID

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14544	44912	В	14630	64	5735	
14545	44913	В	14631	891	6894	
14546	44914	A	14632	33	476	
14547	44915	Α	14633	1	360	GQGEPKQLGRHPGVLRPHVVS
						AKTGDAC/IWTQVRNLFPERPK
						VQWKEGCA\LQSGNSQESVTEQ
						GSKVSAYSLSSPLTLSKADYEK
						HKLYACEVTHQGLSSPVTKSFN
	ļ.,	ļ				RGEIIEGEVPPCI
14548	44916	Α	14634	40	508	LPGECHRAGQQGQRLQPQLLG
						LLMLWVPGSGGEIVMTQTPLSS
						SVTLGQPASISCRSSQSLVHSDG
					NTYLTWLQQRPGQPPRPLIYHV	
					FKRFSGVPDRFSGSGAGTDFTL E/ISGRVEA*\DVGVYYCMQG\T	
						QFPCSFGQGTKLETKRTVAAPS
14549	44917	A	14635	3	652	AWWNSETPAQLLFLLLLWLPY
14349	74717	<u> </u> ^	14033		032	TSGEIVLTQAPGTLSLSPGERAT
ì						LSCRASQTIGSTYLAWYQQRPG
			<u> </u>			QAPRFLIYGASSRATGIPDSSSSS
						SSSSSSSSSSSSSSSSSSSQQ
						YNNWPPGGTFGP\GTKVEIKRT
		İ				VGCTHLFFIFPAHLMSQLKIWE
						LPSACVPVCNNF\YPREVQSYSG
						KVDN/VPSNRVTPRESVTEQD\S
						KDSTLQPSAATLDA
14550	44918	A	14636	3	440	
14551	44919	Α	14637	457	1270	SRRVSFPPSA*QAVSDPSRQQLP
			•			SLRLRWGPEGRSFPAENLGRGV
						RWVGGRDPLAA\LP*SSGPLNA
		1				GRLFARPQESGAPRIGNVRDVP
						RAREDVGTQAGRREREFPRLFG
						PAAVVGCERHLSPFPRAAGAG
						GRAGAAGRRSAPPGGLGSCRSC RISAGPHGNDVPTLGGRGW/PL
						*NSGAEKL*EQ*G*/TGTGSWSL
						AAILGGVPREP\GFRGAGRETLG
						PRGWLVMLSRAVLLHSVLLCT
						CHSSWVLLHSDLACHADNNPD
						IQSCLFPMSKY
14552	44920	С	14638	90	465	<u> </u>
14553	44921	A	14639	116	439	
14554	44922	С	14640	278	364	-

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	l e	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14555	44923	Α	14641	116	687	EIKDAALQLPLWRAQDRERAM
		l				KERREKRREREKERDREKKRK
						EKMFCKRSCHTIMEPTPELSCV
						LPCLAPIVLAPPPLPL\PVLPLPY
		1				ATR*LPQQNRQSTLTEAHGL\TR
						GPITLATNRCMPSTTRDRRRLT
						PPSTLPNTTNLSWGVREGPNQD
İ						PEPSLSPCVPHDPFDYNAESLFS
ļ					}	YWLVSVTPMTAHTP
14556	44924	Α	14642	474	1166	HPNRQGLRAPATAPGHPGREQ
						ALCEAPQGPGQKCGPPRAQEA
						Q/GTEPRRASRAAGACRHHWP
						QVA\RPGAGAPAGAARLGGV*E
						PGRAA/APRLPLPSQCCHR*GTS
				:		PAHLSASRSSSATCPWRRTAGR
						SGPSPCMTL\KQRQGHPRDGWI
						RAPSCDVTNRGLSSLLNLGGSA
						QQLLLASRPLGSGSGALEALEH
						GKSRAPWEPWLHGTHNPVHGK
						DPAGSTSHWPRPPAPRT
14557	44925	Α	14643	283	806	PRGSFSSTGRTRNAAGSTSNCSL
ļ						RSLSARIEERKVSRVCGVPNHR
						HLRLGALTTTRPSWNGAIPTLA
						KLPAKRC*RRSGRRSKTWSFPG
						ST*LGNFAEEMICAASARPAGV
						PGPAL\PTPLRAPLPGVPWTSSR
						GRMCARLSCC/HAGRPVPARPG
		<u> </u>				AARARRAAAGERHRPPALRV
14558	44926	Α	14644	1	338	
14559	44927	Α	14645	11	1091	MSNDGRSRNRDRRYDEVPSDL
						PYQDTTIRTHPTLHDSERAVSA
						DPLPPPPLPLQPPFGPDFYSSDT
						EEPAIAPDLKPVRRFVPDSWKN
			1			FFRGKKKDPEWDKPVSDIRYIS
						DGVECSPPASPARPNHRSPLNS
				Ì		CKDPYGGSEGTFSSRKEADAVF
	1					PRDPYGSLDRHTQTVRTYSEKV
						EEYNLRYSYMKSWAGLLRILG
						VVELLLGAGVFACVTAYIHKDS
						EWYNLFGYSQPYGMGGVGGL
						GSMYGGYYYTGPKTPFVLVVA
						GLAWITTIIILVLGMSMYYRTIL
						LDSNWWPLTEFGINVALFILYM
						AAAISLM*MIPTRGGLCYYPLF
						NTPVNA\GSAG*KEDR*LQ*SSC
		1	L			LSP**FISLVLWFA

RCPFGGLRH*SAR*QPPSGLR GARPGGEWLPC*FPP*PLN WKLPPRLVT*GEPWAAGEKI RLGAGTGPAAGEGLPRARDW WKLPPRLVT*GEPWAAGEKI RLGAGTGPAAGEGLPRARDW PGRAGPERWAPGRGPAGTPATAVKRSAPLPGSGDYANG KPPPSPFCQVXSGNL*VLC4* SLPAA*KPKKA*RRGRRSGYSGP TGYQVQQTRPKNDPKSSGP KNRQSLCLQPLWSFLIQ GRFAALFRASRGERLSLLKLG QLPLPFRALSQGYKPLIGILLA FQRCTAQRGGI*RGSLATVTI CGGLCPV*TSQRRCL;PVRGK TQASVMANAPPRTRLQHLRS* DCCAGSENFKPVDLSLLGSV GSTELDHLAPWLQPAFQGSE FYLTF* LVC4* LLRGVSQ*GYMGVRDPLEEL* CPLAELEHCAGRSAALFRAG* ERLSLLKLRPQLPLSPSA* L4930 A 14648 169 487 EVCYYPPSETYFCQFVKLILC* LFPCWQGVVIQPKLSCMRCL* ULLRGVSQ*GYMGVRDPLEEL* CPLAELEHCAGRSAALFRAG* ERLSLLKLRPQLPLSPSA* L4932 B 14650 I 1683 14564 44932 B 14650 I 1683 14564 44932 B 14651 I 529 MDSEVQADDEVSDENGELIC* WGKGGMGVSDPLEEAVCPL* LERSAGRSAALFRAGSCL* LKLT*/LTAAPSPRGTAFSSFP* PAQRRSIC*Q*Q*LYSCKTFST* PMPRLCL*PQASGKVQ*CEN*S** IPA*QGTSPMCSLCSGAPHQP* APLVLPLPPPVNRLRRELQCV** LARCL*	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
14560	NO:		hod	1	1		
14560		sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
RCPFGGLRH*SAR*QPPSGLR GARPPGGEWLCPC* FPPPLN WKLPPRLVT*GEPWADGEKI RLGAGTGPAAGEGLPRARD\ WKLPPRLVT*GEPWADGEKI RLGAGTGPAAGEGLPRARD\ PGRAGPERWAPGGPAGTYMG TAAVKRSAPLPGS\GDYAMG KPPPSPPCQVKSGNL*VL.C4* SLPA*KPKKA*RRGRQSPC TYQYQVQTPRNDPKSS\GDYAMG KPPSPPPCQVKSGNL*VL.C4* SLPA*KPKKA*RRGRQSPC KNRQSLCLQPLWSFLIQ GRFAALFFASRQERLSLLUG QPLPPRALSQGYKPLIGUL_AFQRCTAQRGGI*RGSLATVTI CGGLCPV*TSQRCLLPVRGK TQASVMANAPRTRLQHLRS DCCAGSENFRVDLSLLGSV GSTELDHLAPWLQPATQGSE FYLTF 14562					sequence		
GARPPGGEWLCPC*FEPPPILL WKLPPRLVT*GEPWAAGEKI RLGAGTGPAAGEGLEPRARDN PGRAGPERWAPGROPAGTPATAVKRSAPLPGS:GDYANG KPPPSPFCQVKSGNL*VLCG* SLPAA*KPKKA*RRGRQSPC TGYQVQTRPKNDPKSSiGPS KNRQSLCLQPLWSFLIQ KNRQSLCLQPLWSFLIQ GRAALFRASRGERSLSLKL QLPPRALSQGYKPLIGLL/ FQRCTAQRGGI*RGSLATVTI CGGLCPY*TSQRCLPVTKG TQASVMANAPPRTRLQHLRS DCCAGSENFRPVDLSLLGSV GSTELDHLAPWLQPAFQGSE FYLTF 14562	14560	44928	Α	14646	398	1107	SPGLVCGLLIFCKVTVPICAGPY
WKLPPRLVT*GEPWAAGEKI RLGAGTGPAAGEGLPRARDV PGRAGPERWAPGRGPAGTPP TAAVKRSAPLPGS\GDYANG SIPAA*PKKA*RRGRQSPC TGYQVQQTRPKNDPKSS\GPS KNRQSLCLQPLWSFLIQ MGVRDPLEAVVCLALELQHI GRFAALFRASRQERLSLLKLI QLPLPRALSQGYKPLIGLLIA FQRCTAQRGGI*RGSLATVTI CGGLCPV*TSQRCLLPVRGK TQASVMANAPPRTRLQHLRS DCCAGSENFKPVDLSLIGSV GSTELDHLAPWLQPAFQGSE FYLTF EVCYYPPSETYFCQFVKLILC LFPCWQGVVIQPKLSCMRCL LLRGVSQ*GYMGVRDFLEEL LLRGVSQ*GYMGVRDFLEEL LLRGVSQ*GYMGVRDFLEEL LLRGVSQ*GYMGVRDFLEEL LLRGVSQ*GYMGVRDFLEEL LKLT/LTAAPSPRGTARSRQECL LKLT/LTAAPSPRGTARSRQECL LERSAGRSAALFRASRQECL LKLT/LTAAPSPRGTARSRQECL LKLT/LTAAPSPRGTAFSSPF PAQRSIIC*Q/LYSCKTFSTT MPRLCLPQASGKVQ*EN*S IPA*QGTSPMCSLCSGAPHQE APLVPLPPPVNRLRRELQCV LARCLPK WLPSFVFETQGPCI STQGTLLVYGLQRPWEKHSI R*HPPSOHGPSQL.PLARGGSS TPCASRMRQHPT 14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARGS SPRLPFCRSAYQPAAGAGGG							RCPFGGLRH*SAR*QPPSGLREN
RLGAGTGPAAGEGLPRARD\ PGRAGPERWAPGROPAGTPA							GARPPGGEWLCPC*FPP*PLNSR
PGRAGPERWAPGRGPAGTPA							WKLPPRLVT*GEPWAAGEKFP
PGRAGPERWAPGRGPAGTPA							RLGAGTGPAAGEGLPRARDWE
TAAVKRSAPLPGSIGDYANG KPPPSPFCQVKSGRL*VLCSLSLPAA*KPKKA*RGRRGSP* SLPAA*KPKKA*RGRRGSP* TGYQVQQTRPKNDPKSSIGP* KNRQSLCLQPLWSFLIQ GRFAALFRASRQERLSLLKLG QLPLPPRALSQGYKPLIGILL* FQRCTAQRGGI*RGSLATVTI CGGLCPV*TSQRRCLIPVRGK TQASVMANAPPRTRLQHLRS* DCCAGSENFKPVDLSLLGSV GSTELDHLAPWLQPAFQGSE FYLTF							PGRAGPERWAPGRGPAGTPAK
KPPPSPFCQVKSGNL*VLCG* SL.PAA*KPKA*RRGRRQSGP TGYQVQTRFKKA*RRGRRQSGP TGYQVQTRFKKA*RRGRRQSGP KNRQSLCLQPLWSFLIQ KNRQSLCLQPLWSFLIQ GRFAALFRASRQERLSLLKLG QLPLPFRALSQGYRPLIGHLA* GRFAALFRASRQERLSLLKLG QLPLPPRALSQGYRPLIGHLA* GRFAALFRASRQERLSLKLKLG QLPLPPRALSQGYRPLIGHLA* GRFAALFRASRQERLSLKLKLG QLPLPPRALSQGYRPLIGHLA* GRFAALFRASRGELSGAVTTT CCGGLCPV*TSQRRCLPVRGK TQASVMANAPPRTRLQHLRS* DCCAGSENFKPVDLSLLGSV* GSTELDHLAPWLQPAFQGSE FYLTF* 14562							TAAVKRSAPLPGS\GDYANGER
SLPAA*KPKKA*RRGRQSPC TGYQVQQTRPKNDPKSSIGP KNRQSLCLQPLWSFLIQ 14561 44929 A 14647 I 540 MGVRDPLEEAVCALAELQHE GRFAALFRASRQERLSLLKLL QLPLPPRALSQGYPLIGNLL FQRCTAQRGGI*RGSLATVTI CGGLCPV*TSQRRCLIPVRGK TQASVMANAPPRTRLQHLRS DCCAGSENFRPVDISLLGSV GSTELDHLAPWLQPAFQGSE FYLTF 14562 44930 A 14648 169 487 EVCYYPPSETYFCQFVKLILC LFPCWQGVVIQPKLSCMRCL LLRGVSQ*GYMGVRDPLEC CPLAELEHCAGRSAALFRAG ERLSLLKLRPQLPLSPSA 14563 44931 A 14649 I 210 14564 44932 B 14650 I 1683 14565 44933 A 14651 I 529 MDSEVQADDEVSDENGELIC LERSAGRSAALFRASRQECLS LKLT/LTAAPSPRGTAFRSPFF PAQRRSIC*Q/LYSCKTFRST* PMPRI.CLPQASGKVQ*EN*SS IPA*QGTSPMCSLCSGAPHQP APLVLPLPPPVNRLRRELQCV 14566 44934 C 14652 295 430 14567 44935 A 14653 452 637 14568 44936 A 14654 151 446 MVLSHWHSRCHWGMKKNY LARCLPKWLPSFVFETQCPC STQGTLLVYGLORPWEKHSI R*HHPSQHGPSQLPLARGGSS TPCASRMRQHPT 14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARS APRSFTPPSPSPSPTQLGLSG GPESAPTA/PGNPSWRSSRWC SPCAASST*KSPYP*CSPT/CA SPRLPFCRSAYQPAAGAGRG							
TGYQVQQTRPKNDPKSSiGPS KNRQSLCLQPLWSFLIQ							,
14561 44929 A 14647 1 540 MGVRDPLEEAVCALAELQHI		1					-
14561					İ		
GRFAALFRASRQERLSLLKLG QLPLPPRALSQYXPLIGULX	14561	44929	A	14647	1	540	
QLPLPPRALSQGYKPLIGVLL/FORCTAQRGGI*RGSLATVTI	. ,						-
FQRCTAQRGGI*RGSLATVTI CGGLCPV*TSQRRCLIPVRGK TQASVMANAPPRTRLQHLRS DCCAGSENFKPVDLSLLGSV- GSTELDHLAPWLQPAFQGSE FYLTF							_
CGGLCPV*TSQRRCL\PVRGK TQASVMANAPPRTRLQHLRS DCCAGSENFKPVDLSLLGSY GSTELDHLAPWLQPAFQGSE FYLTF							
TQASVMANAPPRTRLQHLRS DCCAGSENFKPVDLSLLGSV. GSTELDHLAPWLQPAFQGSE FYLTE		1					-
DCCAGSENFKPVDLSLLGSVG GSTELDHLAPWLQPAFQGSE FYLTF							,
GSTELDHLAPWLQPAFQGSE FYLTF		1					
14562							
14562 44930							, , ,
LFPCWGGVVIQPKLSCMRCL NLRGVSQ*GYMGVRDPLEE. CPLAELEHCAGRSAALFRAG ERLSLLKLRPQLPLSPSA 14563	14562	44930	A	14648	169	487	
14563 44931 A 14649 1 210 1683 14564 44932 B 14650 1 1529 MDSEVQADDEVSDENGELIC WGKGGMGVSDPLEEAVCPL LERSAGRSAALFRASRQECLS LKLT/LTAAPSPRGTAFRSPFF PAQRRSIIC*Q/LYSCKTFRST*PMPRLCLPQASGKVQ*EN*SI IPA*QGTSPMCSLCSGAPHQP APLVLPLPPPVNRLRRELQCV 14566 44934 C 14652 295 430 295 430 295 44935 A 14653 452 637 44936 A 14654 151 446 MVLSHWHSRCHWGMKKNY LARCLPKWLPSFVFETQGPCI STQGTLLVYGLQRPWEKHSI*R*HHPSQHGPSQLPLARGGSST*PCASRMRQHPT 14569 44937 A 14655 2 335 EDRSAFRRQPHTLHPLHARS APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSSRWC SPCAASST*KSPYP*/CSPT/CASPRLPFCRSAYQPAAGAGRGS	14302	14730		17070		107	-
CPLAELEHCAGRSAALFRAG ERLSLLKLRPQLPLSPSA 14563 44931 A 14649 1 210 14564 44932 B 14650 1 1683 14565 44933 A 14651 1 529 MDSEVQADDEVSDENGELIC WGKGGMGVSDPLEEAVCPL LERSAGRSAALFRASRQECLS LKLT/LTAAPSPRGTAFRSPFF PAQRRSIIC*Q/LYSCKTFRST* PMPRLCLPQASGKVQ*EN*SI IPA*QGTSPMCSLCSGAPHQP APLVLPLPPPVNRLRRELQCV 14566 44934 C 14652 295 430 14567 44935 A 14653 452 637 14568 44936 A 14654 151 446 MVLSHWHSRCHWGMKKNY LARCLPK WLPSFVFETQGPCI STQGTLLVYGLQRPWEKHSI R*HHPSQHGPSQLPLARGGSS TPCASRMRQHPT 14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARS APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSSRWC SPCAASST*KSPYP*/CSPT/CA SPRLPFCRSAYQPAAGAGRGS							1
ERLSLLKLRPQLPLSPSA 14563 44931 A 14649 1 210							· .
14563 44931 A 14649 1 210 14564 44932 B 14650 1 1683 14565 44933 A 14651 1 529 MDSEVQADDEVSDENGELIC WGKGGMGVSDPLEEAVCPL LERSAGRSAALFRASRQECLS LKLT/LTAAPSPRGTAFRSPFE PAQRRSIIC*Q/LYSCKTFRST* 9 PMPRLCLPQASGKVQ*EN*SV IPA*QGTSPMCSLCSGAPHQP APLVLPLPPPVNRLRRELQCV APLVLPLPPPVNRLRRELQCV 14566 44934 C 14652 295 430 14567 44935 A 14653 452 637 14568 44936 A 14654 151 446 MVLSHWHSRCHWGMKKNY LARCLPKWLPSFVFETQGPCI STQGTLLVYGLQRPWEKHSI* 14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARS APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSRWG SPCAASST*KSPYP*/CSPT/CASPCASST*CASPCASST*KSPYP*/CSPT/CASPCASST*CASP							-
14564 44932 B 14650 1 1683 14565 1 1529 MDSEVQADDEVSDENGELIC WGKGGMGVSDPLEEAVCPL LERSAGRSAALFRASRQECLS LKLT/LTAAPSPRGTAFRSPFF PAQRRSIIC*Q/LYSCKTFRST*PMPRLCLPQASGKVQ*EN*SI IPA*QGTSPMCSLCSGAPHQP APLVLPLPPPVNRLRELQCV 14566 44934 C 14652 295 430 44935 A 14653 452 637 44654 151 446 MVLSHWHSRCHWGMKKNY LARCLPK WLPSFVFETQGPCI STQGTLLVYGLQRPWEKHSI*R*HHPSQHGPSQLPLARGGSSTPCASRMRQHPT 14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARS APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSRWG SPCAASST*KSPYP*/CSPT/CASPRLPFCRSAYQPAAGAGRGS	14563	44931	A	14649	1	210	EKESSEKEK ÇELESI SIK
14565 44933 A 14651 1 529 MDSEVQADDEVSDENGELIOWGKGGMGVSDPLEEAVCPL LERSAGRSAALFRASRQECLSUKLT/LTAAPSPRGTAFRSPFK PAQRRSIIC*Q/LYSCKTFRST*PMPRLCLPQASGKVQ*EN*SVIPA*QGTSPMCSLCSGAPHQPAPLVLPLPPPVNRLRRELQCV 14566 44934 C 14652 295 430 14567 44935 A 14653 452 637 14568 44936 A 14654 151 446 MVLSHWHSRCHWGMKKNY LARCLPK WLPSFVFETQGPCI STQGTLLVYGLQRPWEKHSI*R*HHPSQHGPSQLPLARGGSSTPCASRMQHPT 14569 44937 A 14655 2 335 EDRSAFRPQPHTLHPLHARSAPRE APRSPTPPSPPSPDTQLGLSGFGPESAPTA/PGNPSWRSRWCSPCAASST*KSPYP*/CSPT/CASPRLPFCRSAYQPAAGAGRG					1		
WGKGGMGVSDPLEEAVCPL	14565		A		1		MDSEVQADDEVSDENGELIGN
LERSAGRSAALFRASRQECLS							WGKGGMGVSDPLEEAVCPLAE
LKLT/LTAAPSPRGTAFRSPFK PAQRRSIIC*Q/LYSCKTFRST' PMPRLCLPQASGKVQ*EN*SV IPA*QGTSPMCSLCSGAPHQP APLVLPLPPPVNRLRRELQCV 14566							LERSAGRSAALFRASRQECLSL
PMPRLCLPQASGKVQ*EN*SV IPA*QGTSPMCSLCSGAPHQP APLVLPLPPPVNRLRRELQCV 14566							LKLT/LTAAPSPRGTAFRSPFKS
PMPRLCLPQASGKVQ*EN*SV IPA*QGTSPMCSLCSGAPHQP APLVLPLPPPVNRLRRELQCV 14566							
IPA*QGTSPMCSLCSGAPHQP							PMPRLCLPQASGKVQ*EN*SWA
APLVLPLPPPVNRLRRELQCV							` `
14566 44934 C 14652 295 430 14567 44935 A 14653 452 637 14568 44936 A 14654 151 446 MVLSHWHSRCHWGMKKNY LARCLPK WLPSFVFETQGPCI STQGTLLVYGLQRPWEKHSI R*HHPSQHGPSQLPLARGGSS TPCASRMRQHPT 14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARS APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSSR WC SPCAASST*KSPYP*/CSPT/CASPT/CASPT/CASPT/CASPT/PGRSAYQPAAGAGRGS							,
14568 44936 A 14654 151 446 MVLSHWHSRCHWGMKKNY LARCLPK WLPSFVFETQGPCI STQGTLLVYGLQRPWEKHSI R*HHPSQHGPSQLPLARGGSS TPCASRMRQHPT 14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARS APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSSR WC SPCAASST*KSPYP*/CSPT/CASPRLPFCRSAYQPAAGAGRGS	14566	44934	С	14652	295	430	
LARCLPK WLPSFVFETQGPCI STQGTLLVYGLQRPWEKHSI R*HHPSQHGPSQLPLARGGSS TPCASRMRQHPT 14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARS APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSSRWC SPCAASST*KSPYP*/CSPT/CA SPRLPFCRSAYQPAAGAGRG	14567	44935	Α	14653	452	637	
STQGTLLVYGLQRPWEKHSI R*HHPSQHGPSQLPLARGGSS TPCASRMRQHPT 14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARS APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSSRWC SPCAASST*KSPYP*/CSPT/CA SPRLPFCRSAYQPAAGAGRGS	14568	44936	Α	14654	151	446	MVLSHWHSRCHWGMKKNYLQ
R*HHPSQHGPSQLPLARGGSS TPCASRMRQHPT 14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARS APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSSRWC SPCAASST*KSPYP*/CSPT/CA SPRLPFCRSAYQPAAGAGRGS		1					LARCLPKWLPSFVFETQGPCDV
TPCASRMRQHPT 14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARS APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSSRWC SPCAASST*KSPYP*/CSPT/CA SPRLPFCRSAYQPAAGAGRG							STQGTLLVYGLQRPWEKHSI*A
14569 44937 A 14655 2 335 EDRSAFRPRQPHTLHPLHARS APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSSRWC SPCAASST*KSPYP*/CSPT/CA SPRLPFCRSAYQPAAGAGRG							R*HHPSQHGPSQLPLARGGSSP
APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSSRWC SPCAASST*KSPYP*/CSPT/CA SPRLPFCRSAYQPAAGAGRG					-	•	TPCASRMRQHPT
APRSPTPPSPPSPDTQLGLSGF GPESAPTA/PGNPSWRSSRWC SPCAASST*KSPYP*/CSPT/CA SPRLPFCRSAYQPAAGAGRG	14569	44937	A	14655	2	335	EDRSAFRPRQPHTLHPLHARSL
GPESAPTA/PGNPSWRSSRWC SPCAASST*KSPYP*/CSPT/CA SPRLPFCRSAYQPAAGAGRG					1		APRSPTPPSPPSPDTQLGLSGPTS
SPCAASST*KSPYP*/CSPT/CA SPRLPFCRSAYQPAAGAGRG							GPESAPTA/PGNPSWRSSRWGSS
SPRLPFCRSAYQPAAGAGRG					1		SPCAASST*KSPYP*/CSPT/CAFP
14570 44938 B 14656 373 502	14570	44938	В	14656	373	502	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	1	<u> </u>	1.465	<u> </u>	1400	CODADA A COSED A VIVIDA CO
14571	44939	A	14657	1	400	GSPAPAAMDEERALYIVRAGE AGAIERVLRDYSDKVKSPGPGR
			}			AVSHPRAALRACTSHPSPPPSFR GCAL\PPPKTFPAWACVFVPPAF
						SIHAS/SHPTTPTVAPTPLIPGPA
						A*PPVLLSSVQVTPPESVDVHL
14572	44940	A	14658	659	1177	ARKGSALLRLYLSNAWCVCVK
		-				VLRIFSTASAYITRRDSSLGPVW
						TFLNFSFGSPGEIPTPFFFFPVFP
						LPLVAWFCPASFCLPQCP\LVPS
		Ì				EPSPPPHSGDAHCSPPKTFPAW
						ACVFVPPAFSIHAS/SHPTTPTV
						APTPLIPGPAA*PPVLLSSVQVT
		l				PPESVDVHLCPVPHS
14573	44941	В	14659	1	318	
14574	44942	Α	14660	2	131	
14575	44943	Α	14661	354	762	CPEIRPPS*THGTK/SHFPPTASG
						THSVLEAL/PDTQRQPSPRVHQ
						DPSPAI/PALLPSPHEHGPQHLL
		İ				QLPSHRCHPDTAPFNCDAVKAC
		l				PGHATAALEHPYLSGRQSPPSP
			İ			THLPKLPAGRGYTVCALPKGPG
						ERSAV
14576	44944	Α	14662	157	413	ALVCS/SSLAIREMQIKTTMRYH
		ŀ				LTPVRMAIIKKSGNNRCWRGC
		ŀ				GEIGTLLHCWLDCKLVQPLWK
		ļ				SVW*FLRNLELEIPFDPAIPLL
14577	44945	Α	14663	1926	2448	DFIAVITTRKQLKTKYNAHHSK
						QAITKHFSPKISDLILCLKKKIM
		ŀ				NRHFSKEDIYAAKKHMKKCSP
1						SLAIREMQIKTTMRYHLTPVRM
						AIIKKSGNNRCRRGCGEIGTLLH
		ŀ				CWWDCKLVQPLCKSMWRFLR
						DL\DPAIPLLGIYPKDYKSCCYK
14570	14046	ļ	14664	1	1504	DTCTRMFIAALFTIAKTWNQP
14578	44946	A	14664 14665	1	1584 4729	
14579 14580	44947 44948	A	14666	1	1566	
14581	44948	A	14667	1	1424	
14582	44949	A	14668	1	576	
14583	44951	A	14669	1	3325	-
14584	44952	A	14670	1	2195	
14585	44953	A	14671	1	3162	
14586	44954	В	14672	287	2018	
14587	44955	A	14673	1	1073	
14307	17733	1^	1,4073	1.	11073	<u> </u>

SEQ ID	1	1	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14588	44956	A	14674	11	3335	MVKGSIQQEELTILNIYAPNTG
14300	44930	^	14074		3333	APRFIKQVLSDLQRDLDSHTLI
	,					MGDFNTPLSTLDRSTRQKVNK
				:		DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCKRTEIITNYLSDHS
		l				AIKLELRIKNLTQSRSTTWKLN
		ŀ	}			NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFIALNVYKRKQERSKIDTL
		ı				TSQLKELEKQEQTHSKASRRQE
	•					ITKIRAELKEIETQ
14589	44957	A	14675	1	1838	MIIPIDAEKAFDKIQQPFMLKTL
1 1307	11757	l' `	11075		1.030	NKLGIHGMYLKIIRAIYDKPTA
						NIILNGQKLEAFPLKTGTRQGCP
		ŀ				LPPLLFNIVLEILARAIRQEKEIK
						GIQLGKEEVKLSLFADDMIIYLE
	l					NPIVSAQNLLKLISNFSKVSGYK
						INVQKSQAFLYTNNRQTESQILS
		ļ	İ			ELPFTIASKRIKYLGIQLTRDVK
						DLFKENYKPLLNEIKEDTNKW
						KNIPCLSIGKINIMKMAILPKVIY
						RFNAIPIKLPMTFFTELEKTTLK
						FIWNQKRARIAKTILSQKNKDG
		1				GITLPDFKLYYKATVTKTAWY
						QYQNRDIDQWNRTEPSEIIPHV
						YNHLIFDKPDKNKKWGKDSLF
						NKWCWENWLAICGKLKLDPFL
						TPYTKINSRWIKDLNVRPKTIKT
						LEENLGNTIQDIGMGKDFMSKI
			Ì			PKAMATKAKIDKWNLIELKSFC
						TAKETTISVNRQPTEWEKIFAIC
						LSDKGLISRIYKELKQRHKKKT
						NNPIKKWAKDMNRHFSKEDIY
						AANRHMKKCSSSLAIREMPIKT
1						TMRYHLTPVRMAIIKKSGNNRC
						WRGCGEIGTLSHCWWDCNLVQ
						PLWKAVWRFLKDLELEIPFDPA
1						ISLLG/IYPKDYKSCCYKDTCTQ
						STFTCRQHLPL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14590	44958	A	14676	ī	2354	MGDFNTLLSALDRSTRQKVNK
						DIQELNSALHQADLIDIYRTLHP
						KSTEYTFFSAPHRTYSKIDHLV
						GSKALLRKCKRTEIITNCLSDHS
						AIKLELRIKKLTQNHSTTWQLN
						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWYVKDTRI
			·			SGMLWYVKAVCRGKFIALNAH
						KRKQERSKIDTLTSQLKELEKQ
						EQTHSKTSRRQEITKIRAELKEI
						ETQKTLQKINECRSWFFEKINKI
						DRPLARLIKKKREKNQIDAIKN
						DKGDIITDPTEIQTTIREYYKHL
						YGNKLENLEEMDKFLDTYTLP
						RLNQEEVESLNRPITGSEIEAIIN
						SLPTKKSPGPDGFTAEFYQRYK
						EELRIKYLGIQLTRDVKDLFKE
						NYKPLLNEIKEDTNKWKNIPCS
						WVGRINIMKMAILPKVIYRFNA
]		IPIKLPMTFFTELEKTTLKFIWN
						QKRARIAKSILSQKNKAGGITLP
						DFKLYYKATVTKTAWYWYQN
						RDTDQWNRTEPSEIMLHIYNHL
ľ						IFDKPDKNKQWGKDSLFNKWC
						WENWLAICRKLKLDPFLIPYTK
		1				INSRWIKDLNVRPKTIKSLEEDL
						GNTIQDIGMGKDFMSKTPKAM
						ATKAKIDKWDLIKLKSFCTAKE
						TTIRVNRQPTEWEKFFAIYSSDK
						GLISRRYNELKQIYKKKTNSPIK
ł						KWTKDMNRHFSKEDIYAAKRH
				<u> </u>		MKKCSSSLAIREMQIKTTMRYH
14591	44959	Α	14677	1	3144	
14592	44960	A	14678	1	2745	
14593	44961	Α	14679	1	2742	
14594	44962	Α	14680	1	3264	
14595	44963	Α	14681	1	2982	
14596	44964	Α	14682	1	3102	
14597	44965	Α	14683	1	1443	
14598	44966	Α	14684	1	3723	
14599	44967	Α	14685	2	1841	
14600	44968	Α	14686	572	744	DLLTTIRMPTDVKKCSSSLAIRE
						MQIKTT\M*YHLTPVRMPIIKKS GNNRCWRGCGEI
14601	44969	A	14687	1	3588	
14602	44970	A	14688	1	5569	
14603	44971	A	14689	1	3992	
005	11.273	<u></u>	1	<u> </u>	1	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ŀ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14604	44972	Α	14690	755	2415	QELRVSGWSPASAPANT*CPPA
						QPCPTRRSGGQRAPARTRSRPG
						SRPRTASALPRPAAQKAASRSA
						RASRTNPYTAPKPLPSVESALQ
						EAAIGEAHAQQRVSCTCAAPA
						APPSPAWPWSAARSAAPPPRT*
						ASAPA*SPL/PVRLKKSPGSSNC
						EARGEGWKT/PALPVQLGAHGP
						PQPAAVPCASPGSGAGPRWSRS
						ARRTPRGPLQRGRTWVSPDPGR
						EATPSGTQWPQKTRGSLPGS*C
			<u> </u>			APETPCSAA*SSGTSPPRGAGGC
						CPGTVPPALEGHN*PQRWGSHS
						AGRVE\TRAEGPQCQQGAPAGC
						SLGVGGQRGPPGLPPIPVPGLC*
						TCRSS*MEAATGSVEKCKRMA
						SLMQASRWSFLRVAPVSGQVK
						YTPGTARWAEPSSLGSDTTLVP
						GAS/RAPADGGFGLKFLSHLVS
						DGVLRCGGPRPGTASAPAGLR
						AAAA\PSVPSLVGPWRGGLGAG
		1				WLGFSLWAFSPTDSLSF*PGPA
		1				LLNLMSALNS*KNFFMVAARS
						CGGGTRGRGGTGPSRGRGRRR
		l				PATGRK*KKKKKKKKFTSPPEV
		<u> </u>				PPSPGLPRDNRL
14605	44973	В	14691	1	822	
14606	44974	Α	14692		2789	MAAVRKQLPRGVLGYGGSGAL
1		l				LSTAAADTSLLLGSLQERKSAS
		İ				AGPGYLRSSVAAWSGLWLTGA
		j				SEAHSVVLACFSRKDVEASLSV
						GLPVHPDASCARRLDAADGVA
		İ				GGSPRLPQSTDGWRLTAHSLPQ
						STDGWRLTAHSLPPPTDGWRL
1						TAHSLPQSTDGWRLTAHSLPPP
		1				TDGWRLTTHSLPQSTDGWRLT
						AHSLPQSTDGWRLTAHSLPQST
						DGWRLTIHSLPQSTDAWRLTA
14607	14075	_	14602	176	140	HSLPQPMDGWSCRSAQWCV
14607	44975	A	14693	176	440	
14608	44976	A	14694	114	253	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14609	44977	Α	14695	124	890	GSRAWRDPRLEVAAATTTLFY
	1.277					GYGFVSAELKKASKRMTCHKR
						YKIQK\KAPEHHRKIRGEAKKR
		İ				GHNKPRKDPGVPNSAP\FKEAL
						L\REAELRKQRLEELKQQQKLD
						RQKELEKKRKLET*SWIFKPSN
						V\DPMEK\EFGLCKTE\NKA\RSG
						KQNSKKLYCQELKKVIEASDV
						VLEVLDARDPLGCRCPQV\EEAI
						V\QSGQKKLVL/IYLNKSDL\VP
		l				KEDL\ESWLKLF*RKDLPNSGCF
]				RASTKTQRVKGKITKVSFIS
14610	44978	A	14696	553	997	EVTACWQSSQPSLTLGASSAW
14010	7770	``	14070	333		APTLAALEEPFSPPLHCGSPFLG
						WPRLEPAPSAPPWAPVRPEPPQ
,						QGPPSAPGRPIPSTTQGLRSADA
						GRRTGTQLHRQPAKVCSF/IS*S
		l				QRDHEPTRRNEQLQTRRLKSCN
			<u> </u>			THREDLOLHSRRKKL
14611	44979	Α	14697	1410	2036	TIREDEQEIISRRRE
14612	44980	A	14698	3	630	RYKGSPRPHQTQEPSWLHLVDP
14012	44900	Α .	14076	3	030	APGLQVELPASPAPCARIRQPL
		İ				GGRWDWAPWSRGWCSSGRLG
						QHRSPWSGWEAQAWRAAGPK
						PCPAGRQLRPGEKSSTAPVGRH
						CWGIQYTLRSHWPGC*VPHSSG
		1				PAGLAGCSGTPAGPQAPHAAP
						VPARA/CSLHTSLQAEGVGSSL
İ						GQPRKGLPQCRWGLKGSSNAA
						KVGAQAGEVQARALRTAS
14613	44981	Α	14699	190	383	ASHPLRGLRSGRSLPDERHRLL
14013	44901	Α .	14077	190	1363	HSAQSHRPPKG*GVRAQGAGL
						, ,
14614	44982	_	14700	1	1095	T/WQLHL*PQCGIHWVKPAGLL
14614 14615	44982	A A	14700	1	777	
14616	44984	A	14701		933	
			14703	1	996	
14617	44985	В	14704	978	1207	ASI DI DCI I CODSI DEED DDI LII
14618	44986	Α	14/04	710	1207	ASLPLRGLLCCPSLPEERRPLLH GAQSHRPPKG*GVRAHGAGLA/
	:					`
						WQLHLRPRCGIHWVKPAGLLS LGPHQQSSKSA
14610	44097	_	14705	111	558	LUTTUŲSSKSA
14619	44987	A	14705	1111	228	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14620	44988	A	14706		1103	MKPWALKVSVTALKVARLEFV PPDVRMCSEFLPCGVKLQTFAV SVTALKAARLELFVSPGGLVVS LASGVKLQTFPVSVTAHKSSVD PRSEQQQDLLQRAKEQTFHTVE GDRSGSPPDSGAQLASPSGSRT RAAGGAACQSRALRPHSSALG WSMGLGAVEQGAVLIREARAA HEPMERVGGSGMAGCRSRALP REKAAKARRETEGSAAVSGWL SGIAQWPKIDHEGFRAVKMPQ AAWLLKAVTFIDLTTLSGDDTS
						SNIQRLCYKAKYPIREDLLKAL NMHDKVAAGFPAGQTHLKTRL EEIRLAVEDGATEIDVVINRSLV LTGQWEG/LYSPVVYSALPCNI* HSRLFLPSGNALLLWLWCLHS
14621	44989	Α	14707	2	2300	
14622	44990	A	14700	576	1058	ARSNIRMVTAFRSLPAGSSGAA RA\MSAHNRGTELDLSWISKIQ VNHPAVL\RRAEQIQARRT\VK KDWQAAWLLKAVTFIDLTTLS GDDTSFNI*RLCYKAGYPIREDL LKALNMHDKGIT\TAAVCV\YP ARV\CDAVKGTSRPAG/CVNIPV GISRAAGF\PAG\QTH\LKTRLEE I\RLAVEDGSLQKIDV\VINRSLG A*QAQWE\ALYDEIRQF\RKAC GEA\HLKTILA\TGELGTLTNVY KASMIAMMAGSDF\IKTSTGKE TVNATFPVAIVMLRAIRDFFWK TGNKIGFIPAGGIRSAKDSLAW LSLVKEELGDEWLKPELFRIGA STLLSDIERQIYHHVTGRYAAY HDLPMS
14623	44991	A	14709	576	998	VLQFLKAACPEFVPSGVPMCSE YLPSGGFVVSLASGVKLQTFPV SVTALNALRLELFVPPGGLMVS LASG\GSCRSSR*VLQLIKAVWT QRPLGGRWDWAPWSRGRRSFG EARAAQKPMEGVGGSGMAGC RSRSPAPREGS
14624	44992	A	14710	2	589	RHHPRHAVQARPFPAGPLAGA QRGSGAE*RGPQGS\PGPAGLS GGESL/PGETDRGADKEERGRQ RWQRALGRLSAGADPLPAELR GRRQRRGR*AGQRAAQEAACE APIQGSGARNGRRG*GRRL*GC YQ*V*FPGLTRGWGRGSRPSAV HCGWEPP*AGKPSGQTPRHSG* PAG\VDGLPPKVTGPPPGTPQPR

	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14625	44993	Α	14711	1	1596	MRLTRCQAALAAAITLNLLVLF
						YVSWLQHQPRNSRARGPRRAS
					i e	AAGPRVSLPPHPLTCVWRCSRP
	:					AMDRPPAASRPETTVATEFVAL
						VLDGARAEAHALLERMVEALR
						AVSARLVAAPVATANPARCLA
	•					LNVSLREWTARYGAAPAAPRC
						DALDGDAVVLLRARDLFNLSA
						PLARPVGTSLFLQTALRGWAV
						QLLDLTFAAARQPPLATAHAR
						WKAEREGRARRSALLRALGIRI
						VSWEGGRLEWFACNKKNQRC
14626	44994	Α	14712	225	980	VPSRTWAEYRTRFVPHSLGQPR
						SSHIPQSPQTDQTAQGQQALAP
						PAGRSGQ*GRPGRRARLGPGER
						HS*APPPLSQLLLQTFPHLQALP
						SAHARPSGQPHADRGHSRSPSG
					}	/DLSTSSVLKVVRPQAGGPKQA
						RCPQAGRVCHCPRGGRRGPE/Q
						Q/GG*RPEAPQASPEKGSTGLRP
						SAKDSKNQGTGIFLRCQVAPRP
						EALCCPPPRGADTPVSPCLWVA
						GAGIPLQPPEGAEGAVLGSLCT
						LCPGSPDSPPTS
	44995	В	14713	405	893	
	44996	Α	14714	178	499	
	44997	Α	14715	231	436	
	44998	A	14716	141	393	
	44999	C	14717	97	351	
	45000	A	14718	119	358	 PPCTSLRPLHAFSGKMTHLNRG
14633	45001	Α	14719	21	614	
						TLSLLDSQLHNLIALQMTCFKD VEIPNFFWEPSVTPSHRNINMYF
						PAAVFGFLPISGTLFSYCKIVSSI
						LRVSSSGGKYK/AFTTCGSHLS
						VVC*FYGTGFGGYLSSDVSSSP
						RKAAVASVMYTVITSMLNPFIY
						SLRNRDIKGVLRQPHGSTVQFQ
J		l				` ` `
			1		1	YLLICSIPFVVWVKKGSKVK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	, , ,
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14635	45003	A	14721	696	1791	VWIALKSENITCELLLFKMQTQ
						CQVLVYLFPIFQPSFLGKRCPIS
						TDPQNLTDVSIFLLLEVSGDPEL
						QPSPCWAVPCPCAWVTVLGNL
						LIILAISPDSHLHTPMYFFLSNLS
						LA\DIGFTSTTVPK\MIVDI\QSHS
ĺ						RVISYAGCLTQMSLFAIFGGME
						E\DMLLSVMAYDWFVAICHPL
						YHSAIMNLCSVGFLVLLSFFFF\
			:			LSLSDSQLHNLIALQMTCFKDV
						EIPNFFWEPSQLSHLACCDTFTN
						NIIMYFPTAIFGFLPISGTLFSYY
						KIVSSILRVSSSGGKYKAFSTCG
į						SHLSVVC*FYGRGVGGYLSSDD
						VSSSPRKGAVASVMYTVVTPM
				:		LNPFIYSLRNRDIKSVLRRPTEH
						CLISKK
14636	45004	Α	14722	1	1971	MTRKVTLLLRTWRGGCGRLA
1					,	MGMEATPQGSLHPLFQEHWIR
						AEDFGVLSPMHPNSVQGVDDM
						IRLGDLNEAGMVHNLLIRYQQ
						HKIYVSLPSPVSTGEPLTGANLL
						DREKKMMHRGHCPIALTEGVS
		•				GPDNPEGRARAKAKAKARAEL
						PLPGAEGTDHLLPSQTYTGSILV
1						AVNPFQVLPLYTLEQVQLYYSR HMGELPPHVFAIANNCYFSMK
						RNKRDQCCIISPQNRVGAQTDT
						VLKVFQHLLISLICKTTLQACCI
						PSHPYLSSAPGPPPLTTQCRSYS
						DSLLGPQLFGDVCYNCSHVIEG
						DGEALCSLPHPTCSPGAWVLQL
						VSPLNKAWCVSCFSCSTCNSKL
						TLKNKFVEFDMKPVCKRCYEK
						FPLELKKRLKKLSELTSRKAQP
		1				KATDLNSAEGPLAQLGAVGA/V
						TGITHEPHIPQVENGSQQLDDV
						PALGV\ENWRIFMAERMEAQG
						GQRLLLSTMHEEYEFVSPSSVAI
						AELVALFLEGLKERSIFAMALQ
						DRKATEGGPVGLTKKQGLLAS
						ENWTLGQNDRTGKTGLVPWH
						CLYTIHTVTKHSDSCCTAVPTL
						LEASSGCIFMSYKKHITLTPKSA
						DSNKRPSTLGSQFKQSLDQLMK
						ILTNCQPYFIRCIKPNEYKKPLG
						FSDSFMLIKLIRIDKVQYDSFSYI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14637	45005	Α	14723	1	1197	MNMNIKKIVKQATVLTFTTALL
14057	13003	 	11723	1		AGGATQAFAKENNQKAYKETY
						GVSHITRHDMLQIPKQQQNEKY
-						QVPQFDQSTIKNIESAKGLDVW
						DSWPLQNADGTVAEYNGYHV
						VFALAGSRKTLMTHQSTCPKH
			ŀ			RASRVGPSNSLQNKQTVKKGD
						QTSVRRKRNDLKVEERHKNHH
						LARELTSSAAAASPTGHAAATA
				:		QAPQRHVEVGRAAAPAGDLGE
						GIVGEQVLGAERLVGPDASRAR
						ALCVLGDRREAGHTAEMTAAP
						PAQVPGPGGPTALRRPSSYLGL
						GVDEQNPMSVDGDEDDVRVQ
						HQELRDCLIKHTILIRGEFVTRS
						LNIAQAADRRDAFVKGIYGALF
						LWIVKKINAAIFTPPAQDPKNV
				1		RRAI/GLLDIFGFENFENNRIIRT
11620	15006	_	1.470.4	ļ	1000	LCWRYREARC
14638	45006	C	14724	1	1032	METTELEOOLNI I LEHEHKOKO
14639	45007	Α	14725	1	4960	MSTTSIFQQLNLLLSHSHKQKQ
						QIQLPNGRSRYSCLMAAFRMQ
						YFLSLPMKSLILVGFERLQAIAR
1						SQPLARQYQAMRQRTVQLQAL
1						CRGYLVRQQVQAKRRAVVVIQ
						AHGQGMAARRNFQQRKANAP
						LVIPAEGQKSQGALPAKKRRSI
						YDTVTDTEMVEKVFGFLPAMI
						GGQEGQASPHFEDLESKTQKLL
						EVDLDTVPMAEEPEEDVDGLA
1						EYTFPKFAVTYFQKSASHTHIR
		L				RPLRYPLLYHEDDTDCLAA
14640	45008	Α	14726	86	370	RTWMTHVWRRRRVQAAPTAY
						SSFRPGTSPCPAQPRAGARA*TR
		į				SGTSIMPTGFSSHTSCLTVP\PFT
				!		ASSAGQHRQRHRKTTAAGHPT
						TTGAASF
14641	45009	Α	14727	3	455	MQSVLPAHGVSSTRASSRSVTL
						SSPSTSTNCRGRKSSRVT\SPVG
						SGPGGGGGGPGSGRFGIDPPRK
						FAAAGSFSTGAPSSVMVVRVSG
						MPSLSW*QYP*SCSTLGCRSRSS
						STISLSSLWA/VQPIAAPPYPELA
						TQLPISCR*SPSPCAPS
L	J				<u> </u>	

SEQ ID	_		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	1	09/540,217	codon for peptide sequence	or peptide sequence	detection,possible nucleonide insertion)
14642	45010	Α	14728	338	1782	NQLSK*KAHSIRT*TALQQSSGR
1						SQICIVQSSADALPSSRCTNPSST
						FTPTPLPPRPHPRTCSTLGCRSR
						SSSTISLSSLWAASR*QPLRTPN
						SRTQLPRSWPLKSTALSPPPPGQ
		1	ŀ			ATRLCWWMVSSK*RTWMTHV
		l			:	WRRRRVQAAPTAYSSFRPGTSP
1		1	•			CPAQPRAGARA*TRSGTSIMPT
			•			GFSSHTSCLTVPRLHCVLCWTA
						SSTPQESSPGGWTRPWPSWRRC
						CTCTGAGRYLQNSTAS/T/PVPS
		1				TSVSQTAPSPKEDYRCWPSYHH
						GSLPPFSVQPG*GCGMSVRAMP
		1				SVGPLWSPTRPPGQIKRQNYLL
				İ		NYWSCWKMTLLAGIIPSST/GS
		l				RPADLDYFLGSSQCPRDERTLK
						GHVSCDPTSSISITGEKLKQKLE
				:		S/SSTLADTKSAFLQDSWT/SSLP
		1		† !		TLGASAI\RTTSAS*TDQCPPAP
						GPLHVSF/SLNSTYPSRLCSSTCS
		1				TVDSFQIIPMPN*MPFSTCGAFC
						WCLWFSAITALYFVM
14643	45011	Α	14729	148	1040	WCTGSQFSKGTWEILLRKKPSL
		l				GNVGALPCVATACLTTVEKRA
						PRGHPLAPVLGDICPKVGSMSS
						STSEL/CTWGNFEFWGRNKYNH
						WF*WGWALLHQFKMGKGAQG
						LQEASFYRKDTY*A\TMP*NPIG
						ISEFGT\LALPGPCKNVFERFMS
		1				RFELPGKAAAMTDNTNVNYVR
		1		ļ		YKGDYYLCTETNFMNKVDIET
İ		1				LEKTEKMWKPRFRGICPQLQG
		1				CSVTEQGFEFSDDAKSWALQH
		l				PKRPAQQDWAMSRGNQGWAN
		1				LGLKIIRGEAVTTRAKEVSVVL
						KVLEVTDGKVWAREKKLRDSR
14644	45012	Α	14730	189	507 -	AAGADTSQGPGGPPPSQSQR\
						PKSPSAGRKGSQLSPQPQKKGL
		ŀ				PSPQGTRKSAPSSKATPQASEPV
						TTQLLGQPPTQEEGSKAQGMR*
						PPQRLGASTQQQTNQEAA
14645	45013	Α	14731	2	356	DGVSLCHPGWSAVARSQLTAT
						SASWAVCWQGCGETGILLPCW
						WECEKVQPLWKQYGRLSCEDH
				İ		LNS*VQVILTTQPPILFPKWLHL
						FTFPPTGQKDSSFSTTLPTHCPG
						GRGCSELRSRHCTPAWVTE
L			L	1	l	

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14646	45014	A	14732	1	408	FFFFFYTTFFLFFYFYFFEMESH SVTQAGVQWYNLSSLQPPPPRF RRFSCLSLPSS*DYRCPPPHPAN FLCF**RQGFTLLAKMVSIS*PH DPPTSASQSAGHVLTSKWELNI EYTWVIKMGTADTGEHKNGEG WE
14647	45015	C	14733	262	536	
14648	45016	A	14734	179	745	TDCLHNRCHRSPRSPPPGPPG*R SRADWWRRGRCWGSWCASSR RVC*RCPAPSASFSCCRSPRGRQ GRTFPPSPCPAPD*TQ\PVSPGDS TRPCPPPSSPPAHRAT/GCCGRR RLPGNASYLASPGKEAGEFPAS QGDS*PGQYGYIGDSSS/RPTRG TWMAKASTPRHPIAPQICGA*G FGEHQEVPVREK
14649	45017	В	14735	1	1164	
14650	45018	A	14736	35	542	VRPPSHVTADSGRSPLSLTYLPL QEPGDMAAAVPRAAFLSPLLPL LLGFLLLSAPHGGSGLHTKGAL PLDTVTFY/KDYGDKLNMELSE KYKLDKESYPVFYLFRDGDFEN PVPYTGAVKVGAIQRWLKGQG V\YLGMPGCLPVYDALPGEF\IR ASGVEARQALLKQGQDN
14651	45019	A	14737		908	VRPPSHVTADSGRSPLSLTYLPL QEPGDMAAAVPRAAFLSPLLPL LLGFLLLSAPHGGSG\LHTKGA LPLDTVTFYKVIPKSKFVLVKF DTQYPYGEKQDEFKRLAENSA SS\DDLL\VAEVGISDYGDKLNM \'ELS\EKYKL\DKESYPVF\YLFR EGDF\EEPSSHYTGGS*RLGAIQ RWLKGGKGVYLGDGLELVCT V*LTPWSGEFIRA\SGVKARQAL LKQGQDNLSSVKETQKKWAEQ YLKIMGKILDQGEDFPASEMTR IARLIEKNKMSDGKKEELQKSL NILTAFQKKG\AEKEEL
14652	45020	A	14738	3	227	
14653	45021	Α	14739	1	424	
14654	45022	Α	14740	1	207	
14655	45023	Α	14741	2	352	
14656	45024	Α	14742	1	441	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14657	45025	A	14743	1	461	MTVDYRKFNQVVTPMAA/AVP DAVSLLEQINTFPGTWYAAIDL ANAFFSIPVHEAHQKQFAF\LPQ GYINFPALCHNLIRRELDFFLLL QDITLVHYIDDILLIGSSEQEVV NTLDLLIHKRSKEAEHTAASRIR VSCLPEQKSHEQTLPWEQVP
14658	45026	Α	14744	1	928	
14659	45027	Α	14745	283	1074	
14660	45028	A	14746	450	1970	VRVLSPVEKELKLWKNTHKLL SYPTVGAAVTQLQNLTAMGVI GSHGARGQVVALNRQRQGDL QPFTRVTVHWGKGYDRTFWGL LDTGSELTLTPGDPKHHCGPPV KVGAYGGQVINGVLAQVQITV GPQTHPVVISPVPECIIGIDILSS WQNPHIGSLTGIMVGKAKWKQ LELPLPRKIVNQKPYCIPGGTVE ISATIKDLKDAGVVIFTTSLFNS PIWPVQKTDGSWRMTVGYRRL NQVVTTPTAAAAPDV/VVSLLE QINTSPGTWYAAIDLANAFFSIP VHKPHQKQFAFSWQGQQYAFT VLPQGYINNPALCHNLIQRNLD HFPLLQDITLVHYLNDIVLIGSA IKWVMHSS/DSIIKWKWHVHDR AQADPEGTS*PGYGHF\WCPIC QQQRPTLSSQYDTIPWAISQLPG DRLIIGPLPLWKGQKFVFTGIDT YSGYGFAYPACNASAKTTIMES QNTLSTLMVFHTALPLTKALTS WLKKCSSGLMLMEFTGLTMFSI

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14661	45029	Α	14747	1	2589	MGPPYKPPTTWPPQTLSYHSTT PVKLQSLSLPPTAARGATEPCH
						LQAPASSVQQMTPPAPGLKHL
						GGKSQMCLEQDGRSGSVRGSP
						CHAGHWVAGAGFGKGTTRTW
						GHVNLIVPPNCPTASSSAARDS
						APNQPLLGVVTAQIVPPTCPPA
						MTVQELIAPLTLPPAAGCRRGV
						RPFQTRVWLLAQLLRARLGGP
						GEGVVTCHSWRLERVAPRLAS
						LWWALAAETAARLDWAAKRP
						SAWGSIKAAEFSSKATPRLEGA
						SSRTLWVSYIQAKETTTGKNGP
						VEPSHLWEAYCRATAPHPAVH
						SPSAEVTYTCGIVNLLLQGLVP
						KSHLETHTGCHVVVELIICSVV
						LPLISRLSDPDWIHLVLMVIFSK
						ARDPTPCPASGPEQPSVPTSLPL
						IAEVQQLPEGRAPSPVAAPGFLS
						KQIQDALCALEGSQALEPKGGE
						GSEGVEAEGPGTETETVLPVST
						LNFCPEIQIDIADKEIEQGEVTTS
						VTALLEGLEKTCSSRPSCLEKD
						LTNDVTYLDPTIFSFEPLSSPDG
						PVVIQNLHITGTITAQELSGTGL
						QPCTRYTVKCETSLDGENSSLQ
						QLAYHTVNHRYWEFLNLQ/SH
						QEKKPDL*KFIKN\IEKMVVSAI
						VDNLKTAFPCSESQGPTEELSE
						AETESKSQTEGKKARKSRLRFS
						SSKISPALSVTEAQDTILYCLQE GNVESKTLSMSGMESFIEKOTK
14662	46020	_	14748	470	2507	
14662	45030	A	14748	478	3507	SKILRAVIWGSHLVRSAMKTET VPPFQETPAGSSCHLNNLLSSR
						KLMAVGVLLGWLLVIHLLVNV
						WLLCLLSALLVVLGGWLGSSL
						AGVASGRLHLERFIPLATCPPCP
						EAERQLEREINRTIQMIIRDFVL
						SWYRSVSQEPAFEEEMEAAMK
						GLVQELRRRMSVMDSHAVAQS
			į			VLTLCGCHLQSYIQAKEATAGK
						NGPVEPSHLWEAYCRATAPHP
						AVHSPSAEVTYTRGVVNLLLQ
		1				GLVPKPHLETRTGRHVV
14663	45031	Α	14749	1	417	
14664	45032	A	14750	34	350	RERGQEGGVAGRVEMDVSVVE
						RGAYADQIWGRPKGGESRGEK
						RKGRWVQRVRALRARVTPAGL
						GECKWGRGRGD*VRRVKPRAL
-		ŀ				RERWGDGSTVSDSTPMKPGLE

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	1 -	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14665	45033	Α	14751	1	1604	MASKRCIRDFRSRGGKLKSFQL
						YATFRKEGCGPVLANGNPGGG
						RGQEVSATQDLRRGWGIASSAS
						RFPWRQSWPGRPRLGNYDSGA
						PGEAKMPSKGKDKKKGKSKGK
						DTKKLIKTDESVVDRAKANASL
						WEARLEVTELSRIKYRDTSRIL
						AKSNEDLKKKQCKMEKDIMSV
						LSYLKKQDQEKDNMIEKLKQQ
						LNETKEKAQEEKDKLEQKYTR
		ŀ	ļ ļ			QINELEGQFHQKAKEIGMIHTE
						LKAVRQFQKRKIQVERELDDEI
		ŀ				NDLLVKEKIMQLVQQRSQIQTL
						QKKVVNLETALSYMTKEFESE
						VLKLQQHAMIENQAGQVEIDK
	1					LQHLLQMKDREMNRVKKLAK
	ł					NILDERTEVERFFLDALHQVKQ
						QILISRKHYKQIAQAAFNLKMR\
						EHVQEEQNIPKSEHLMAESTAPI
	ŀ					V*IRIFWRPKNGHILKEMWILEI
						*PGSRRKKYCDCSLQK*MAVLL
		l				GNTTQSSRPSSSRTMLFLDSGET
		l				KEFGDESKLQDKIFITQQIAISD
						VFW*SGATHYSKRTSGV*HSGK
						SESLQPRGQRLI
14666	45034	Α	14752	95	473	
14667	45035	Α	14753	168	449	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	ļ			sequence		
14668	45036	Α	14754	 1	2000	MAFEQAKQVVKLALDLWPIGD
14000	43030	``	14734	1	2000	GPVELQVTVLDQHANWSLRQE
						QHGKRGPLQFTDVAIEFSLEEW
						HCLDTAQRNLYRNVMLENYRH
						LVFLGIVLSKPELITCLEQGKKP
j						LTVKRHEMIAKPPGQKNWEKQ
						ALPLTGCLTKYPSWAKNIDIIVT
						CSHFARDLWPEQSIKDSFQKVT
						LRRYENYGHDNLQFKKGCESV
						DECKVHKRGYNGLNQYLTTTQ
						SKIFQCDKYVKVIHKFSNSNRH
						KIRHTGKKPFKCIECGKAFNQS
						STLTTHKKIHTGEKPFKCEECG
						KAFNWSSHLTTHKRIHTGEKRY

						KCEDCGKAFSRFSYLTAHKIIHS
						GEKPYKCEECGKAFKRSSNLTT
]					HKIIHTGEKPYKCEECGKAFKR
						SSILTAHKIIHSGEKPYKCEECG
						KAFKHPSVLTTHKRIHTGEKPY
						KCEECGRAFKYFSSLTTHKIIHS
						GEKPYKCEECGKAFNWSSHLT
						THKRIHTGEKPYKCEECG\KGF
						KYSSSLTTHKIIHTGQQPFKCEE
						CGKAFKCFSILTTHKRIHTGEKP
						YKCEECGKAFNSSSHLTAHKRI
						HTGEKPYKCERCGKAFKRSFIL
		İ				TRHKRIHTGEKPYKCEECGKGF
						KCPSTLTTHKTL\EIHAGKKPNK
						*E/ECSKAFKDTSALTRHKIIRTG
						EKPCEFDECRKAFNQLSTFTKY
14669	45037	В	14755	269	1131	
14670	45038	Α	14756	2	149	ALFRAGRQECLSLLKLCPQPPL
						SPTCFVPGR*GFYLHFFQRCAA
				ļ		QRGGI
14671	45039	Α	14757	2	175	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
14672	45040	A	14758	1523	2667	PLLVIHRQTGSGEDLQQTPADL
						QQRDLLERKLTNRNE*HQHQQ
						KRHPLKNPTQMSP/CIKDQRYIN
						PRSTIKLELGIKKLTENCTTTWK
						LNNLLLNDCWEKQERSKTDTL
						TSQLKELEKPEKTNSKASRRQEI
						TKIRAELKEIGTRKTLQKINKCG
						SWFFEKINKIDRSLARLIKKKRE
						KNQIDTIKNDEADITTDPMEIQT
						TNREYYKHLYANKLENLEEMD
						KFLDTYTLPRLNQEEVESLNRPI
						TSSEIEAVINSLPT/KKRSPGSDR
						FTAEFYQRYKEEL\VPFLLKLFQ
						TIEKEGLLPNSFYEDSIILIPKPG
						RDTTKKENFRPISLMNIDAKILN
						KILANRIQQHIKKLIHHNQIDFIP
		1				GMQG\WFNIHKSINVIRHLNR
14673	45041	A	14759	640	1771	APVARMTA/PLFYTALLVFSAL
						GNILALCLTCQ\KRRKINCTGIY
						LVHLAVSDLLFTVALPGRVVC
	}					YVLGSSWPFGKGLCRLTAFVL YTDTYGGVYLMACVSVDHYPT
						\VVCAHWGPCLRTAGRARLVC
						VAIWTLVLLQTMPLLLMPMTK
						PLVGKLACMEYSSMESVLGLPL
						MVLVAFAIGFCGPVGIILSCYM
						KITWKLCSTAGRTQ*PAGKDTT
						GGASPGGPSDQPEKTPPAGAAQ
						EDPVTPRERTPPAGQPERTQ*PA
						GKDATGGTAREDPVTSRKGRH
						WRGCLLTLLMLVAVVVCFSPY
						HLNIKQFMARGMLHLPSCAER
						RAFLLSLQATVALMNMNCGI/D
						PIIYFFASTHYRKWLLGILKLKG
İ						SSSSSSSSTLY
14674	45042	Α	14760	1	401	MVLVQWLTAITSAFWETQNKG
					+	SLQFEDKWDFMRPIVLKLLRQE
						SVTKQQWFDLFSLGATLQDSSG
						PLLTGKLKKKRKVSGTKY/IEAS
						SATSLKATTDTHHLLLYNPFWI
						PLTLI*HLCWLQRFA*SCDTDPH
	ļ	1_				P
14675 14676	45043 45044	A	14761 14762	34	2720 251	
14677	45045	A	14763	2	452	LQIEII*ALEELAAKEKANEDAV
,	13373	l.,		[PVCLAADFPSAGMGSSCDGQE
						DELDIKSRAAYNVTLL/NFMDP
						PKIPHLKEKPYFGMRKMAVRW
						HHAENLVDRPQ\EESEDDSHLE
						GRDLDTWHVGVKISWDVKTPG
						LATPQGDCYFMLGNLWRTKIIL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14678	45046	Α	14764	1	1701	MPFKCKKQHTFNKTSPKEGAII
						FTHGCLCTSPFNSELSEHLGNL
						KLSSWEHLHHRSQQTLSVRLH
·						VPTPQPVEHEFIQLIYIHTGAKL
				:		QASKVIIATKDGKRRKSPLWQL
						KYPKLILREASSVSEELHKEVQ
						EAFLTLHKHGCLFRDLVRIQGK
	ļ					DLLTPVSRILIGNPGCTYKYLNT
						QLFTGP\FPVKGSNIKHTEA*I\A
						AVCETFLKLNDYLQIETIQALEE
						LAAKEKANEDAVPLCMSADFP
						RVGMGSSYNGQDEVDIKSRAA
						YNVTLLNFMDPQKMPYLKEEP
						YFGMGKMAVS\WHHDENLVD
						RIQLQLWDPVDGTTASSAVPKQ
					1	GRDKRREVPNTLSQPNSQEGLI
						SAFHSRCVHAVFYKLTSAPKTT
i						SLPDVPSDLLAEGLAASRAGQA
		:-				LWRCDKKSLLTLVRKINLQGK
						RDGGRSATNQQSQPHLDNTGL
						ERQLEQKVGRCRSPWRGLGHL
						EGNAQKGHLLHGVNKNGQHL
						LRRHGVSIRAGRAPHLAAADT
	j					APGRISCLGHSEALKPRIQAGGL
						GAESIWQVLWEGGSNSSVFCR
						HDEEPAAELPLPGKQASGGHM
14679	45047	В	14765	16	1462	
14680	45048	Α	14766	89	566	MEQGSIITQAWREDALVLTKQ
						GLVSK\SSPKRPHVCHILK\APFC
						CSCTQHVGQSSSPR*ANPHIRRE
						STPIAMSDQLQRLQYQFYQIPG
						TCLLPEVTEKNQGRICMVTDLD
						ETLVHSSFKPISNADCLVPVELE
						GTMHQIHVLMRPYMDEFLT*M
						EEMF
14681	45049	Α	14767	1	305	LSGLEMFQAGAGAPSSPGY*GS
						CWAISWSGPPDKHSNRGASCG
						LETWAGAAATDTAGPHPAAGR
						C*A*AGSGGDQGTGSTAPLPAT
						P*PLQSLV*AAALPA

14682	L		09/540,217	location of first codon for peptide sequence	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14002	45050	A	14768	1070	2653	SLKPCMACL*HS*EKTLSDSSG RSWICAVSF\GSPGPAAAPLPEL SFPLRPGYTKLLPHATEEVSPFG SGAGRKLGPMLPSTAELCPGTR ARWSEGYRDLGRQAQDLQAG RKSVVTWALEGHPREEERAAA KVQQCQRMWEGCSDPQELIPR PGRNPDPVPDSAVALLPWGPSR VPDLGPDSEHR/PAHSRVQYWL HKAPAPPSPSPQLAEPATPPTRP SSPGSRSPSSWHRLRSWSLNAD PW\RPRPADLPPGNPAPPSRGGA HGTWGKDVRPLGRGSRLFSVV CGRREVQRAARRVGTTGPGWR APGREREAASR*QPPGTG/PASP MGRASSSGSRVSPGPIRRTAPW EGCSCRSRHHWGKGLGPPDGE PG\PGSS*VAQPTVGGRRSLGPG GAAPARDAAAATPPGDAAARE GSGRHLRAPASRPQHQPRSPRS HPPASATTASLQKFRSASSEPPS EDRPFDFSPTAPPQCPRVAPSAG AR*R/AGGRLPE/PR/RPGPWGG GSRTNLTIPTSLHGPTPAAARTS PHRKSR
14683	45051	A	14769	748	1073	SQHFGRPRWVDRLRSGVRDQP GQHDKTPSLLEMQK\IAGRAGG CL*SQLLRRLRQENPLNPGGGG CSELRSGLCTSVWATGRDSI*K KKKKKKKRKPWSLLSLIVLVQF
14684	45052	В	14770	59	466	
14685	45053	Α	14771	2	3184	HRTGIPGSTISSPGGETVPE\MTA AMRERFDRFLHEKNCMSDLLA KLEAKTGVNRSFIAL\GVIGLVA LVLVV\GYG\ASLPLQPLLGFCY PA\YISIKAIGE/SPNKEDDTQW/ V*TYWVV\YGVFS\IAE\FFSDI\F LSW\FPF*LHG*KCGFLVGWCN GPEAPSKWGLKLL*QAASSGPF LSLEATKSQIGQVLVKD\LKDQ GPKETADAIH*KKPKKSLP*ILL G*KKKKK
14686	45055	A	14772	328	479	PQFPCRRFRALSEEELDNEDYY SLLNVRREASSEELKAAYRRLC MLYHPDKHRVPELKSQAERLF NLVHQAYE\VVERRRTPAEIRE EFERLQREREERRLQQRTNPKG TISVGVDATDLFDRYDEEYEDV SGSSFPQIEINKMHISQ

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14688	45056	A	14774	15	1095	KVAKMATALSEEELDNEDYYS
						LLNVRREASSEELKAAYRRLC
]	ļ					MLYHPDKHRDPELKSQAERLF
						NLVHQAYEVLSDPQTRAIYDIY
						GKRGLEMEGWEVVERRRTPA\
						EIREEFERLLREREERRLQQRTN
						PKGTISVGVDATDLFDRYDEEY
						EDVSGSSFPQIEINKMHISQSIEA
						PLTATDTAILSGSLSTQNGNGG
						GSINFALRRVTSAKGWGELEFG
		ŀ				AGDLQGPLFGLKLFRNLTPRCF
						VTTNCALQFSSRGIRPGLTTVL
						ARNLDKNTVGYLQWHCSSPLL
						QVQRPHRNTRACAPEPSFRPFL
						HVPTWDAECSGARTPSTAWTS
						AAVKLREACLSGPGSGSHQLLL
<u> </u>						LTPRSKRRTGGG
14689	45057	В	14775	508	1765	
14690	45058	Α	14776	2	639	GRRRHRRRHRDCSRGDCYLPT
						GTNTQQPLPPPREPLPELPARTP
						DLSMQNSEGGADSPASVALRPS
						AAAPPVPASPQRVLVQAASSNP
						KGAQMQPISLPRVQQVPQQVQ
						PVQHVYPAQVQYVEGGDAVY
						TNGAIRTAYTYNPEPQMYAPSS
				,		TASYFEAPGGAQVTVAAS/SPP
						AVPFHSMVGITMDVGGSPIVFS
						AGAYLIHGGMDRPELLANL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14691	45059	Α	14777	1	2091	MKPDDNSTQNEGGQSFEERHE
						TLQRVKFRGGPKTAPGRKLIAF
						LGQMNSGTSGGQRVPGEEELM
				ľ		AGGATCLPNTEAHQPIRMKNL
						HTPAFRLCPLTLQDNEADAFW
		1				HQAFVDWGMPTGKKPQTLGLT
İ						YKTTHDLSMQNS/EGGADSPAS
ļ		l				VALRPSAA/APPVPASPQMLPLE
		l				LVILGVV/PLDKCIPAQWSKEG
]						QTNSYTYNPEPQMYAPSST/AS
						YFEAPGGAQ\MVGITMDV/GGS
		1				PIVSSAGAYLIHGG/MDSTRHSL
		l				AHTSRSSPARSFRSNSKAPSSRK
]						FCC*LPTLSSKTL/SLFSKNLLRK
				1		LPLGKHSPDSVQWPWIYGLLV
]				AGSESLASLLFPKASSSGETLGF
		1				ATGQLAGESRLPSLLAFILVPID
		1				VAWGKEQLVTQTTVYRVQGS
		1				GQHLNKAWQEPALVNICGMGE
						KVDVQVGLCQHRAHLDGWNS
						SRAASVWPGETLTHHQDQCQA
		1				NLPPGIETSILGTKGARYPSPQE
						QGGGRVWLERLDPRG/AVSGT
						DL*CLPQSDVPPGRA*SGGTGD
						KSISTHDKSGRGNSKYHY/YGIR
						LKPDSPLNRLQED/TQYMAMR
		l				QQPMHQKPRWCRAELGGAGSP
						L\GMEDVSHVFPEFPAPD/LGSF
						LLQDGVTLHDVKAL/QLVYRR
						HC/EDVV/MNLQFHYIEKLWLS
						FWN/SKASSSDGPTSLPASDEDP
	ļ.,					EGAVLP/KDKLISLCQCDPILRW
14692	45060	Α	14778	94	3006	RTRSLTRKAMAEHAPRRCCLG
						WDFSTQQVKVVAVDAELNVFY
						EESVHFDRDLPEFGHVLDVHG
						VHVHKDGLTVTSPVLMWVQA
						LDIILEKMKASGFEFSQVLALSG
<u> </u>						AGQQHGSIYWKAGAQQALTSL
					<u> </u>	SPDLRLHQQLQDCFSISDCPVW
						MDSSTTAQCRQLEAAVGGAQA
						LSCLTGSRAYEFNLVCDRKHLK
						DTTQSVFMAGLLVGTLMFGPL
						CDRIGRKATILAQLLLFTLIGLA
				<u>L</u>		TAFVPSFELYMALRFA\GL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14693	45061	A	14779		710	MGPAAGAPTSRASRGPRAPALR AAVGRRCPRPEMPCQCHPLAR CGLDPAWTQLPQETAGAPRQD PRCTNQLICEHLPLEGLGFGYL GNKDSLRHVKVVAVDAELNVF YEESVHFDRDLPEFGGLASSAV TQSSS/DDLQQVP*SLQSLHAA* SLQTNASPFTFEHSFLGKRHPK GAGQKRAESPTRPPLRLQEAAP GELQRPLPYGTQVPAGLCPSVS GLRLRARALLPYRTHSLSV
14694	45062	В	14780	1	252	
14695	45063	Α	14781	73	198	
14696	45064	A	14782	762	1080	AGRAEMQQERESHEEAKDPSP LSPREGSQKQQQNVNTSTPSHC PR/PPLRMPPPAPRRPVP/CDHPR AEECERKAQHWQAAPPAAPVR DPLGEASWAPESGGDVESLYI
14697	45065	A	14783		309	MTKTEREQNEQLDGGGYGERL GEYSLFLLCVPSELNQGKCERV SSYFGMNLSGVLKKLPFPPTRE SWPPPHFTGTPHNHHGGVSNG ALGCAQRCSLHHATNAGGWER GGVSNSRLFFFLLSASFSDAKIK PAPSPIDHLRAEEC*/PPPLSQPP ALVAWCREHLWAHPKAPLLTP
14698	45066	В	14784	1	471	
14699	45067	В	14785	1	1143	
14700	45068	A	14786	1992	2205	WRSMSGLRKRQLASQISYCPC WS\CP*EESHCRRVSPAPLA*LL SPCPAPQQRSHTDPPPFPFKHFP LPFSG
14701	45069	В	14787	389	973	
14702	45070	A	14788	287	525	TNPFNLLQH*KKP*CYTYISITD PTDACHKNKAFSSPAVSQLT/CI FSSTAGSFFVLSGHSCGHLAVFS LWPRSPQLPGTR
14703	45071	Α	14789	475	616	
14704	45072	A	14790		704	MTKWKSNPQIHMELQRTLSNP NNLEKSNVKTYYKAKIFDDNLI QLTFQQHRQDRDTEQEKKKEE EREKRKQQQEPLRKKAKMQTV RPGFDLTQGPNCLTPHGKTWSR LPPKNHRRPYTSRPEGPRVKA* NHGPEGRRGTRLPPEAAKRAQ AEEAAQ*WPTRSTLRLEVTPRA MPP\SPSEDPRVAGVHPAPPRGP RSDS/APLRPQLRPPRSLFSLASE PAAAGNARGRHSVLALY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	•	
14705	45073	Α	14791	200	408	SPHLMSINLITFANFLFPCE*RNS
						QFLKS/SIFVMWMIHNGMKCRQ
						LDTCHCRHGTCKDSICPGLEES
						PLP
14706	45074	Α	14792	240	771	GRGMQSPAQVDPAAKRGNIAG
						HTAHLWPKGGTLPLCF/YHLLI
						MQIREQP*SSLEYPTF/RILSLVS
		l				GTKLNKILANRIRQHIKKLIHHN
			,			QVGFIPGMQGGLPTPGSEQGHL
						RVSPSSPSSTVTLKLFNFMEPFE
						NLNLSKGKASVHTITQENTDSL
						KVQGSQFSQGHEKRTLDLKVS
14707	45075	В	14793	1	1557	
14708	45076	Α	14794	150	1668	ESDARQGPFPLENTRPHPSCMQ
						NHGICRSLHPGLCGCPPTARDA
		İ				APATAGGVPAY/GPLTGAPPG*
						LPGAGIGSKQSSTMGPSGAGSC
						LGL*APSQPAD/PGSIQGVLVIA
						DTE*PLVGGDGGLEAEHQWQ*
						EAGP/GGLIPKHMNPQSRQQSLF
						GRRGYRDSSLRSVLGRTGQQT*
						HAELGAGVMGPGGNGR*GH/G
						PRGLPCLVQGRRGPGVAAVAA
						GVSVAVPSGMLSENGPGCSK/P
						HSGDAGQRRGLRENL*GVRALI
]					WGAASSFLLTEQDLQELADCL
						HSGRKTCFFYFQGRGDVAPGPV
		•				APAHHRLAGISTVVPEGPGLCV
					:	GEDRQLPG/GNQRAEPPP*LVRS
						VSSLTGLPALGPRQMPNAGAG
						NGSGK*RGVRGSGPDWDCTGT
	1	ŀ				HGGAGTGKLLWAGLSLFPLSG
						KSEAHQQRAAGHGYGRGLGSP
						GRPPRSRKGRVLGSESAPA*GSP
		l				QPLRSARQKPESSCPRQGQPVG
						RGPGAPLEAGHCSFPRPLGPAR
		<u> </u>				SLHPDFPQGLQA
14709	45077	Α	14795	3	374	RRPPSWG*RGCSQLPECAADGS
						LEDHGPLPPMVSCFSRHPAGRF
						LRVFPPSPLLAVDCCWHNNSYV
						VAVDCCSATSCSLLLYPSQEFPP
						HKFLVYLVLSWHQLLGGLELIC
		<u></u>				TQQPVNPVVSDG

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14710	45078	A	14796	73	890	MSLVRTDGKVSFPHSRAASSPP PRFAGSLPWTPKGHGFTSFPRL EGYCPS/REGPSDWGRNSFRVSS WAWRPSQGVSHHTPGQGP/PSP GNRDK*VATRKSPGQPGSSPEP GWSLQSTSSPGATRTKGNLNTL KG*KQVVD*WVGVGTGWEGS GAPQSPQPSREWRLSPRGVGVV KETWPSPWSGR*APPCPGQPQG QQTHMERGGR*G*QVGACGLG P*GQWSSVTNGKDTRAPWSPPP RQRNGRWPAPPPFTFHYYYYY YYYYYYYFFFFGE
14711	45079	A	14797	273	678	TEESTSVLSVPNQTGAGAARIGI VSLRLCGLYVTSPPRAARMHIR PHGPWPGGSHHENGEAGPES/E GAPASASGRGAPEGQAWPPRD AVLSSLMLALFRTKSARHLVLG TRQTAASPFPFFSQPSWYCSLG VDGG
14712	45080	Α	14798	551	743	PTRITRSAGSSSWGSCRPRAPAP FSWSRPCLQGPRG*CRGLHSRF YAAP*VQGHPVHILPYSKF\RRG RAALRPRGQWGLDSVLGTAVC GCGP/RSESDICSGR*GPE/WYIS PPVPPHVHRGFQQL\NCARQVG KGGGGPGCPMLGNH\SSQVGL GLGGQDQDSLESPGRVSSPSGP SAASDPGG*GFQGNPDPVHPAP APPDW

SEQ ID			SEQ ID NO:	1		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14713	45081	I A	14799	[] [705	2299	KREWRPRHHPLGPPPRCAGRSA
14713	45001	^	14///	1,03	LLJJ	PLSGPASPFS*WLPPGHGPCGR
						MCSRAARGGDVTYRTHNTRAP
						SCPSLHWVGERG*DAHTRTPSC
						PSLHWVGERG*DAHTRTPSCPS
						LHWVGERG*/GCPHSGLQLSFP
						PRSAGCCKALFEAEKSKGDTGL
						GKRGWAGGGAGGAARPCICNP
ľ						TVPCPQSTQSPPV/PPGTR*QLG
ŀ						SSASHSRAASSPPPRFAGSLPWT
						PKGHGFTSFPR/RGGVLPFPEGP
]		SDWGRNSFR\GPAGPSMW*PAP
						PSDAQVGHGPGRAGLQRCRGR
						GGHTPFVLAAAAIDSAYFS**/D
						P*VFGSNFPHLSCFPG\GCGEGN
						Q*ARRRPLAGAAALPGPNLES
						WVCELSVAGPAVWDPVPSP/GD
					}	PTTGPESP/RNSAGACGAQGSM
						R*T*GTPATWSL/GPPPELPGSR
						HQRQCECGPAHVQRERMPGW
	•					ACMLKSVAQRGFQRR*QLSKA
						QAAAWRV\PGGR\GAGKD*GW
						RRWGVSRGGGGLCCARGGGGS
						SVRRGEVEGS/GCARGGGGCAG
						WRALSPSSLPISGAQELGS
14714	45082	A	14800	1	267	CSAGGPWRAPQPRRFHRRRP
						AQLPPPLPLPPLPASPRIHNRTF\
						PRPSQRTPPPAALGCPEPGS/RS
				İ		QGRGHARPPGSGEGDPTVSSPG
14715	45083	A	14801	218	430	
14716	45084	Α	14802	9	201	
14717	45085	Α	14803	1453	1936	EVEKHLCQG*ELLRAQHN*AA
						ACRRPRPPAPGPQCSAGGPMAR
						APAPQVPPPPPPCSA\PPPLPLPP
						LPAS/HAHPQPHFRHGRRSALLP
						RPPWAVRSRGALAGPRTRAAA
						GLRGGAGAAPAPADARFPASSP
						AE*PKFPQNSARALTGFPRCTD
						PTVSSPGY
14718	45086	Α	14804	1	580	TSGCSHSLHKPAGVSTRPPTVA
						SICAPRPRTSPLRRGSLTLWPFP
						GGHLTLLETP\GFDSCPEEVEGP
						RFSTFKTS*ARGGRRQRCAPRG
						RREVGPPSAAESGRARGEPPRC
]						RLLCRGGK/DPQPRAGEGEAAA
]						GTPEGSRAGASGSEGALPSAAS
						PGNSSGRSRPEAGANPKGAPRR
		L				CVWGEPGGRRTRLFPDE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14719	45087	Α	14805	279	1148	ALHEFAMSRSLHASLH*S**STC
						P*QQPAPAPAGEPRARFPGKSSS
						ASSGPG/WPRNHGKQLHLPAQP
						Q/QQGRISSPRVAGGSGCTGRA
						LLLAAGLQNSTRSLNPRDPPTIL
1						SHRPLQGM/PDAP/RGCSHSLHK
						P/SRS/TPLSGQL*PPSAPHAPAQ
						APSVAGPLPCGLSPGDT*RYWR
		ŀ				LLRFDSCPEEVEGPRFSTFKTS*
	-					ARGGWRQSRVWASARGASTL
						QASVQGR*NPQPRAGEGEAAA
						GTPEGSRAGASGSEGALPSAAS
		Ì				PGNSSGRSRPEAGANPKGAPRR
						FLGGCLL
14720	45088	Α	14806	765	4270	SQDTPGDLQCHGDLL*DGAVLS
						EAPGGERVPRDD*GGVCPEGVP
						HVSLLRDSVAEALQPRGEAGG
						DQVPEAHARPPSAQR*PAGAGL
						RLHPPAAGGVPGQSGGHLPKIQ
						DDLTSKS*TNSITKTLFPNKVTL
						SDSGRA*ILEGHRSTQGSLGLEA
						DPGTVSHHQHPCPPNAATHHF
						HRTARPGVQQGPGPASVQQPES
						DGDGALLRSPCSLLPQGADEVL
						LQPDGDKQGGRPRGDSESD*G
						YSERR*AQDEYQGH
14721	45089	В	14807	133	2943	
14722	45090	Α	14808	1	814	MEEELLRELLYSENWWSIPYND
						TIQELCCWAVLKRTFINGNTSR
		ŀ				KPGMKMVTGLKKQERRSKQW
						GCEKEAKFVPDMWIVLEVGRP
		İ				EERGIDGKRWGKSCSVEQSVW
				ļ		AFQNPIWSKAQTLLNSMK/AYE
		ļ			1	KGKEGVEEKFEARRSWFMRKK
				ļ		AISHLHNMEMQGEAASADVEA
		ļ				AVSYPEDLAKIIDESDYTKQWIS
						NVDKTVFYWKKMPSRTFIATE
		l				EKSMPGFKASKDSLTPLLGTNA
						AGGF*LKPLLTYHSQTSRALKN
		<u> </u>				YAKLTLLMLYKWNK
14723	45091	Α	14809	123	515	PKQSAKTIWTSTARPEDVVTCS
]			FKEEFLPGNTHHDFSIKNRSDPE
						LEVSRPLARPENFPANRSHSLLE
						L/RPCRGRWAAHWTGCSSLLD
1						ASSAPRTDQPSLWRNLPTASLG
L	<u></u>	<u> </u>				GPWSHNQPGIVFQNTYLMKWP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14724	45092	Α	14810	279	1136	VDVDLACVRGLLCSCDEFLSW
						LALNRTVLATPALPSCLSQMAS
		ĺ				SLPGLCARTRRGERADSQRGTC
		l		1	İ	AFFSPARTASQAPLVASSVRKQ
1						RVWAPNPLNCHTLPPSRGSQRG
						GYIGEEGVFGSRVVGIWVPGPN
						GSPWPLGVQR/PECVLLGVGSG
						S/SSHELFSEADFISRLLAPSPSP
			,	}		APAS/MGNAIEEEAPELGPLGRV
				İ		FVDDFHYDYNFIHLHEDLCYGP
						LEGPDLDLAGTGDRTPPLHSRP
						RASLSPSALRCP*ALGF/WSTLA
ļ]				WGSPNSSSRVPEAQPPAPSLAE
14725	45093	Α	14811	1	1597	GGRKAAVGVETRPQGSAQPTW
						LPVHGDTCTTLLLPCRAWWRL
						GGLPWAASVVGAKCVWTPPER
	ŀ			:		GLLMLHKPMPGWSSGYRVVLP
						AVAFFRLRAQVTKLGEEVSLRF
		1				LKREAKLCGFLQKSFLALEKR
						MKASESSRLKLEGSLRGELESR
						WEKLRGLMEERLRALQGQHEE
						SHLLEQCQGLDAAVAQLTKFV
						QQNQASLNRVLLAEEKAWDA
		1				KGRLEESRAGELAAYVQENLE
		1				AAQLAGELARQEMHGELVLLR
14726	45094	Α	14812	150	1625	CLGKAFSCRELNAYLRLPTFFIV
		1		=		FSLRQPSLRMHSRAWVKTCGA
						MPCSCCRIRHCSSVTDMAVRSL
		İ				ALLQLVNRTPLRRRQISAACQR
						MSTFLFFRACTF*PRNRSSSGRH
						/DLTARPRQAILRQNFLK/RAQS
						RNLTAPNSPR**LLVKTYRQTS
						RPVSAAH*CPR*RGRAALAAAC
						HSGSPGARRQTAGPAEGPAR*T
						RRDFL**RAACLRRRWKWPE*L
						*RSRSSSP*SQGSCPPALPAPPRR
						CSAGSRTGPAARWCRAGAGSR
						RHAGSSSA***SRR*GRSASGRR
		-				PRHGGTAGSGRWLWWARCPA
						*AGIPWPSWPCHSSWP/PQSVRV
						VSSG*RASLHSPGLDSSGGGLQ
			}			SFLLVQLPAQGKTSSQP\RHAA
						APSAAACPHAE*CTLGSCPRSA
						TLSPTPPRASPPAG\RSAVSPCPP
						GHEGVLVISKHHLLLLSRSGRG
						LGGNVVRPCSVQPMDGVGQEL
						AVADRDQLPCLVCQDGECPLK
					!	ASGRPVLGCP
L		<u> </u>	L	<u> </u>		ANGUA FEGU

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14727	45095	IA	14813	921	1446	ACCISHGNLWHHLDQKHGLQS
14/2/	43093	^	17013	1921	11440	*GEKSRLFSRPNLQVPVHTGLD
						HRARYLTGPRKQGHLWPTGPS
						SVFDNENFCKLSVGM*TGRASA
						RQQIQSPFAESGKAFSDICSVSL
						RLCVLPHPWNSRLRSPALRNSL
		1			1	KYSVTSSAAARRKLLPLASGPL
						RLGPRSLPTRLQVLRWEKVASA
14728	45096	A	14814	1	675	REGERSEFIREQUERWERVASA
14729	45097	A	14815	1	2341	MASDLVNRCEACDLPQGPISVH
		` `	1.10.10			QFPRLSNEVTTTVGRNSESPNQ
						RSVSGPWTPREVRRGVAAAAH
						GFSGQNAKWAAHIPATAMSFA
				1		LPYSVLHSPHALTTHPLNPGDA
						VTMPSWTSTVSAANASKGRSFS
						QCFQSTEEGQERRRPRTSQAKV
						QQGSTAGKVNELITLAGHIKSQ
l						APHGKRKPPPRDSHVICSMLRK
						RKPDETPSPSSDAQFSGMLAVA
						VSSLLCPRSSPSQQWHCHSPSTT
						ILGSSDVSDTSFQPSTKLGPVHG
		İ				FPPQSRDCILDFELIRKVTGAWS
						SLTAGSWETKIQEFSEEDILAPH
						RITPGPHSSKSPVHFLASLWGR
						KTKLYHHTIQDKEPRSVPEAGS
						SGRRDGGGAAHSMLCMAPEPD
						HPGSNPNCRSPATLHADADGN
						LQREAAKCQGDGTQVSTHVDH
†						TSAPRQAETGPATGENEKMESH
						GRSKRGLVGADQCPAETLDSY
						TLLVGHKKRDKDRQTPGPAPQ
						VNSFVPENYLLRGPLAPLGSAIS
			•			MKTAGAVHIRPSGDLGRAPAP
						AVRPRPRPPRRPRRLLPGPAHLP
						GAPRPPAPAPLSRRAARQSPAA
						AAAPPPGCE/RRG/PAGPASGAG
]				PPAPAHGARPGGKPAANPPRAP
						QP*P/RRRRSRRRAQPRPGPAPL
				1		SRMRAEGPARALRPPPHYQPEL
				1		GPPSETVGPVTELLAPIPTARFP
14720	45000	<u> </u>	14016	44	455	TRSHPYPVTASPARSS*TSGLD*
14730	45098	Α	14816	44	455	HLRNRTR/PSQITPHIYNHLIFDK
						/PLFNIWWWENWLAICRKLKLD
						PFLTPYTKINSRWIKD*NIRPKSI
1				1		KNLEENLGNTIQHLG/IGKDFM
				1		TKTPKAMATKAKIDIWDLIKLK
						SFCTAKETIIRVNRLPT*WEKIF
L	L	<u></u>	L	<u> </u>		AIYP

SEQ ID NO:	SEQ ID NO: of peptide sequence	ł .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14731	45099	A	14817		279	IQDIGMGKDFMTKTPKAMATK AEIDKWDLIQLKSFCTLKEITIR VNRQTTEN\YPSDKGLISRMLP NF*RLFPAS*KKPPLFDIGKTEK TNYS
14732	45100	В	14818	1	2025	
14733	45101	В	14819	1	3570	
14734	45102	В	14820	1	1380	
14735	45103	A	14821	1391	2742	KKRESSLTHVMRPASF*YQSQA ETQQKKRILDQYP**TLMQKSSI K/YLAKRIQQHIKKLIHHDQVGF IPGMQGWFNIRKSINVIQHINRA KDKNHMIISIDAEKAFDKIQQPF MLKTLNKLELEETTVKFIWNQ KRAHIAKSILSQKNKAGGITLR DFKLYYKATVTKTAWYWYQN RDIDQWNRTEPSEITLHIYNYLI FDKPEKNKQWGTDSLFNKWC WENWLAICRKLKLDPFLTPYTK INSRWIKDLNVRPKTIKTLEENL GITIQDIGMGKDFMSKTPKAMA TKAKIDK WDLIKLKSFCTAKET TIRVNRQPTKWEKIFTTYSSDK GLISRIYNELKQIYKKKTNNPIK KWAKDMNRHFSEEDIYAAKKH MKKCSSSLAIREMQIKTTMRYH LTPVRMAIIKKSGNNRHAPFSIH THIMFGSLYLQIQKDLSILGFW YPRGILEPIY

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14736	45104	Α	14822	1	1698	MEDEMNEMKQEGKFREKRIKR
						NEQSLQEIWDYVKRPNLRLIGV
						PESDGENGTKLENTLQDIIQENF
:		1				PNLARQTNIQIQEIQRMPQRYSS
		1				RRATPRHIIVRFTKVEMKEKMS
						RAAREKEIQTTIREYYKHLYAN
						KLENLEEMDKFLDTCTLPRLNQ
						EEVESLNRTVTGSEIVAIINSLPT
						KKSPGPDGFTAEFYQRQSESQI
						MSEFPFTIASKRIKYLGIQLTRD
		1				VKDLFKENYKPLLNEIKEDTKK
		1				WKNIPCSWVGRINIVKMAILPK
		ŀ				VIYTFNAIPIKLPMTFFTELEKTT
			!			FKFIWNQKRAHVAKSILSQKNK
						AGGITLPDFKLYYKATVTKTA
		1				WYWYQNRDIDQWNRTEPSEIM
		l				LHIYSYLIFDKPEKNKQWGKDS
ļ						LFNKWCWENWLAICRKLKLDP
						FLTPYTKINSRWIKDLNIRPKTIE
						TLEENLGITIQDIGMGKDFMSK
						TSKAMSTKAKIDKWDLIKLKSF
						CTAKETTIKVNRQPTKWEKIFA
						TYSSDKGLISRIYNELKQIYKKK
						\TNNPIKKWVKDMNRHFSKEDI
						YAAKRHMKKCSSSLAIREMQIK
						TTMRYHLTPV
14737	45105	В	14823	91	1596	
14738	45106	Α	14824	1	2757	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		_
14739	45107	Α	14825	1	2677	MGKKQSGKTGNSKKQSASPPP
						KEHSSSPAMEQSWMENDFDEI
						NKIDRLLARLIKKKGEKNQIDT
						TKNDIGDITTNPTEIQTTIREYY
					:	KHLYTNKLENLEEMDKFLDTY
		l				TLPTINQEEIESLNRTITGSEILAI
						INSLPTKKSPGPDGFTAEFYQRY
						KEELVPFLLKLFQSIEKEGILLN
						SFYEASIILIPKPGRDTTKKENFR
						PISLMNIDAKIFNKILANRIQQHI
						KKLIHHDQVGFIPGMQGWFNIR
						KSINVIQHVNRTKDKNRMIISID
						AEKAFDKIQQPFMLKTLNKLGI
						DGTYLKIIRAIYDKHTANIILNG
						QKLEAFPLKTGTRQGCPLSPLL
						FNIVLEVLARAIRQEKEIKGIQL
						GKEEVKLSLFADDMIIYLENPIV
						SAQNLLKLISNFSKVSRYKISVQ
						IPQAFLYTNNRQTESQIMSELPF
						TIASKRIKYLGIQLTRDVKDLFK
				-		ENYKPLLNEIKEDTKKWKNIPC
						SWAGRINIVKMAILPKNWKKT
						TLKFIWNQKRARIAKSILSQKN
						KAGGITLPDFKLYYKATATKTA
ļ						WYWYQNRDLDQWNRTEPSEIT
·						PHIYNYLIFDKPDKNKQWGKDS
					1	LFNKWCWENWLAICRKLKLDP
						FLTPYTKINSRWIKDLNIRPKTI
						KTLEENLGITIQDIGMGKDFMS
						KTPKAMATKAKIDKWDLIKQE
						SFCTAKETTIRVNRQPTKWEKIF
						ATYSSDKGLISRIYSELKQIYKK
14740	45108	В	14826	1	1186	
14741	45109	A	14827	l	2304	
14742	45110	В	14828	1	2995	
14743	45111	Α	14829	1	3095	
14744	45112	Α	14830	1	3654	
14745	45113	Α	14831	1	3189	
14746	45114	Α	14832	1	2064	
14747	45115	В	14833	1	3031	
14748	45116	Α	14834	1	3192	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14749	45117	A	14835	1	3210	MVKGSIQQEELTILNIYAPNTG
14749	43117	l ^A	14633	1	3210	ALRFIKQVLRDLQRDLDSHTIIM
						GDFHTPLSTLDRSTRQKVNKDI
						1
			:			QELNSALHQEDLIDIYRTLHPKS
						TEYTFFSAPHHTYSKIDHIVGSK
						ALLSKCKRTEIITNCLSDHSAIK
						LELRIKNLTQNRSTTWKLNNLL
1						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYQNLWDTFKAVCRGK
						FIALNAHKRKQERSKIDTLTSQL
i .				-		KELEKQEQTHSKASRRQEITKIR
						AELKEIETQ
14750	45118	Α	14836	1	3144	
14751	45119	Α	14837	113	1336	PTLAGTGNQETSHLGFNAIPIKL
		İ				RMTFFTEL/EKTTLKFIWNQKR
			ļ			ARITKSILSQKNKAGGITLPDFK
						LYYKAIVTKTVWYWYQNRDID
						QWNRTEPSEIMPHIYNYLIFDKP
						DKNKKWGKDSLFNKWCWEN
						WLAIYRKLKLDPFLTSYTKINS
ŀ						RWIKDLNVRPKTIKTLEENLGN
						TIQDIGMGKDFMSKTPKAMAT
		İ				KDKIDKWDLIELKSFCTAKETTI
						RVNRQPTEWEKIFATYSSDKGL
						ISRIYNELKQIYKKKTNNPIKKW
			ļ			VKDMNRHFSKEDIYAAKRHMK
						KRSSSLAIREMQIKATMRYHLT
					•	PVRMAIIKKSGNSSRSNIVHSVT
						QANALESSWIPLFLIPHIQYISKP
						QASSFKMYHVTDPILPPSSQHL
						DQAAIINLCLQWPLLWFECMCP
						SKIHMLKLTPQ
14752	45120	В	14838	1	3127	SKITIVEKETI Q
14753	45121	A	14839	1	2742	
14754	45122	A	14840	1	1998	
14755	45123	A	14841	11	3030	
14756	45124	В	14842	11	2558	
14757	45125	A	14843	1	3828	
14758	45126	A	14844	1	4134	MRKKQSRKTGNSKKQSTSPPPK
14730	75120	()	117077		1154	ERSSPAMEOSWTENDFDELRE
						EGFRRSNYSKLQEEIETKGQEV
						ENLEKNLDKCITRITNIEKCLKE
						LMELKAKARELHEECRSLRSRC
						1
		1				DQLEERVSVMEDEMNEMKQE
						GKFREKRIKRNEQSLQEIWDYV
						KRPNLPPIDVPESDRENGTKLE
						NTLQDVIQENFPNLARQANIQI
						QEIQRMPQRYSSRRATPRHIIVR
						FTKVEMKEKMLRAAREKAFKQ
	1					ASRREDDIAKVTSG
14759	45127	В	14845	1	3555	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14760	45128	Α	14846	1	3345	
14761	45129	Α	14847	1	3451	HTKNPSVHHHRQRPKVDKTTK
						MGKKQNRKTGKSKTQSAFPPP
		ŀ				KERSSSPAMEQSWMENDFDEL
						REEGFRRSNYSELREDIQTKGK
						EVENFEKNLEECITRITNTEKCL
						KELMELKTKARELREECRSLRN
						WIKSQDPSLCCIQETHLMCRDT
						HRLKIKGRRKIYQANGKQKKA
						GVAILVSDKTDFKPTKIKRDKE
						GHYIMVKGSIQQEELTILKIYAP
						NTGSPRFMKQVLSDLQRDLDS
						HTLIMGDFNTPLSTLD
14762	45130	Α	14848	1	2606	MVASRSSRVGVFYGGQNSVDG
						VSRRSDGSLNELCLYRPPTGAY
						LAYDTLDVLPSGQFTVNTPPID
						VNGKSLLALLWEHTSLLTSMG
						GGRWAMQPLGQLGQLEHLGD
		ł				RRDQVLDRSGRSRRLDAAQHL
		ļ				RHRPDRRPQGMVGVLTNQKEP
						RTRWIHSRILPEVQGGTGTVPSE
						TIPIDRKEGILPNLFDEASIILIPK
						RGRDTTKKENFRPISLMNIDAKI
						LNKILANRNQQHIKKLIHHDQV
						GFIPGMQGWFNICKSINVIQHIN
						RTKDKNHMIISIDAEKAFDKIQ
						QPFMLKTLNKLGIDGTYLKIIRA
						IYDKPTANIILNGQKLEAFPLKT
						GTRQGCPLSPLLFNIVLEVLAR
						AIRQEKEIKGIQLGKEEVKLSLF
						ADDMIVYLENPIVSAQNLLKLIS
						NFSKVSAYKINVQKSQAFLYTN
						NRQTESQIMSVFPFTIASKRIKY
						LGIQLARNAKDLFKENYKPLLN
		•				EIKEDTKKWKNIPCSWVGRINI
						VKMAILPKVIYRFNAIPIKLPMT
						FFTELEKTTLKFIWNQKRARIA
					}	KSILSQKNKAGGITLPDFKLHY
		1				KITVTKTAWYWYQNRDIDQW
		1				NRTEPSEITPHIYNYLIFDKPEK
		1	1			NKQWGKDSLFNKWCWENWLA
						ICRKLKLDPFLTPYTKIHPRWIK
						DLNVRPKTIKTLEENLGNTIQDI
						GMGKDFMSKTPKAMAAKAKI
	<u> </u>	 				DKWDLIQLKSFCTAKETTIRVN
14763	45131	A	14849	1	2862	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14764	45132	Α	14850	1	3724	MGKKQNRKTGNSKTQSASPPP
İ						KERSSSPATEQSWMENDFDELR
						EEGFRRSNYSELWEDIQTKGKE
						VENFEKNLEECITRITNTEKCLK
						ELMELKTKARELHEECRSLRSR
						CDQLEERVSAMEDEMNEMKRE
						GKFREKRIKRNEQSLQEIWDYV
1						KRPNLRLIGVPESDVENGTKLE
}		:				NTLQDIIQENFPNLARQANVQI
		:				QEIQRMPQRYSSRRATPRHIIVR
						FTKVEMKEKMLRAARQKAPH
						HTYSKIDHIVGSKAL
14765	45133	Α	14851	2	2375	
14766	45134	A	14852	1	2376	MGDFNTPLSTLDRSTRQKVNK
		ŀ				DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIL
						GSKALLSKCKGREIITNCLSDHS
						AIKLELRIKNLTQNRSTTWKLS
						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDTFKAVC
						RGKFIALNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQKTLQKINESR
						SWFFERINKIDRPLTRLIKKKRE
						TNQIDAIKNDKGDITADPTDRQ
ŀ						TTIREYYKHLYANKLENLEEM
						DIFLDTYTLPRLNQEEFESLNRP
ŀ						ITRSEIVAIINSLPTKKSPGPDGF
			İ			TAEFYQRYKEELRIKYLGIQLT
						RDVKDLFKENYKPLLKEIKEDT
		-				NKWKNIPCSWVGRINIVKMAIL
		ŀ				PKVIYRFNAIPIKLPMTFFTELE
						KTTLKFIWNQKRARIAKSILSQ
						KNKAGGITLPDFKLYYKATVT
						KTAWYWYQNRDIDQWYRTEP
						LEITPHIYNYLIFDKPEKNKQW
					ĺ	GKDSLFNLWCWENWLAICRKL
						KLDPFLTPYTKINSRWIKDLNV
						RPKTIKTLEENLGITIQDIGMGK
						DFMSKTPKAMATKDKIDKWDL
	1					IKLKSFCTAKETTIRVNRQPTK
						WEKIFTTYSSDKGLISRIYNELK
						QIYKKKTNNPIKKWAKDMNRH
	1					FSKEDIYAAKKHMKKCSSSLAI
						REMQIKTTMRYHLTPVRMAIIK
L	<u> </u>	<u> </u>	<u> </u>	I	<u> </u>	1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14767	45135	A	14853	1	1576	MDKFLDTYTLPRLNQEEVESLN
11707	1.5.15.5		1.1033			RPITGAEIVAIINSLPTKKSPGPD
		1				GFTAEFYQRLISNFSKVSGYRIN
						VQES\QAFLYTINRQTESQIMSA
		1				LPLTIASKRIKYLGIQLTRDVKD
						LFKENYKPLLNEIKEDTNKWK
						NIPCSWVGRINIVKMAILP/KEL
						EETTLKFIWNQKRARIAKSILSQ
						KNKAGGITLPDFKLYYKATVT
						KTAWYWYQNRDRDQWNRTEP
						SEIMPHIYNYLIFDKPEKNKQW
						GKDSLFNKWCWENWLAICRKL
						KLDPFLTPYTRINSRWIKDLHV
		ŀ	}			RRKTIKTLEENLGNTIQDIGMG
						KDFMSKTPKAMATKAKIDKW
	·	l				DLIKLKSFCTAKETTIRVNRQPT
						KWEKIFATYSSDKGLISRIYNEL
}						KQIYKKKTNNPIKKWAKDMNR
		ļ				HFSKEDIYAAKKHMKKCSSSLA
						IREMQIKTTMRYHLTPVRMAII
						KKSGNN/R/CAPGTPERQNHSLS
			,			WKGS*SQEPSGLAQWIPLPWSP
						AS*DPLA*NSCCQHSCLKSTWD
						AQACARKLRTLIKGYRNCQLE
14768	45136	В	14854	328	1467	
14769	45137	Α	14855	3	3412	
14770	45138	Α	14856	1	2907	MDKFLNTNTLPRLNQEEVDSL
						NRPITGSEIVAIVNSLPTKKSPGP
	į					DGFTAEFYQRYKEELRIKYLGI
						QLTKDVKDLFKENYKPLLKEIK
		1				EDKNKWKNIPCSWVGRINIMK
		1				MAILPKVIYRFNAIPIKLPMTFFI
		į				ELEKTTLKFIWNQKRALIAKSIL
						SQMNRAGGIMLPDFKLYYMAT
						VTKTAWYWYQNRDIDQWNRT
						EPSEITPHIYNYLIFDKPEKNKK
		1				WGKDSLFNKWCWENWLAICR
						KLKLDPFLTPYTKI

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14771	45139	A	14857	1402	3004	TEPKTKTI*LSA*MQKGPLTKF NNPSC*KLLIN/IVLEVLARAIRQ EKEIKGIQVGKEEVKLSLFADD MIVYLENPIVSAQNLLKLISNFS KVSGYKINVQKSQAFLYTNNR QTESQIMSELPFTIASKRIKYLGI QLTRDVKDLFKENYKPLLNEIK EDTNKWKNILCSWVGRINIVK MAILPKVIYRFNAIPIKLPMPFFT ELEETTLKFIWNQKRARIAKAIL SQKKKAGGITLPDFKLYYKATV TKTAWYWYQNTDIEQWNRTEP SEIMLHIYNYLIFDQPEKNKQW GKDSLFNKWCWENWLAICRKL KLDPFLTPYTKINSRWIKDLNV RPKTIKTLEENLGITIQDISMGK DFMSKTPKAKATKAKIDKWDL IKLKSFCTAKETTIRVNRQPTK WEKIFATYSSDKGLISRIYNELK QIYKIKTNPIEKWVKDMNRHF SKEDIYAAKKHMKKCSSSLAIR EMQIKTTMRYHLTAVRMAIIKK SGNNSLIVGASMDTSLLSGPIPY DIOHFDRFGSVTVRRGCAIEFY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14772	45140	A	14858	884	2925	GIKYLGIQLTRDVKDLFKERS/Y
	1				1	EPLLNEIKEDTNKWKNIPCSWV
	:	•				GRINIVKMAILPKVIYRFNAIPIK
	1	İ				LPMTFFTELEKTTLKFIWNQKR
						ALIAKSILSQKNKAGGITLPDFK
						LYYKATVTKTAWYWYQNRDI
						DQWNRTEPSEITLHIYNYLIFDK
						PEKNKQWGKDSLFNKWCWEN
						WLAICRKLKLDPFLTPYTKINSR
						WIKDLNVRPKTIKTLEENLGITI
						QDIGMGKDYMSKTPKAMATK
						AKIDKWDLIKLKSFCTAKETTIR
	•					VNRQPTKWEKIFATYSSDKGLI
						SRIYNELKQIYKKKTNNPIKKW
						VKDMNRHFSKEDIYAAKKHM
						KKCSPSLAIREMQIKTTMRYHL
						TPVRMAIIKKSGNNSPTCVSQD
						WKLTQAVGLWSAEAEAPKSAQ
						RRPLAEGHRAAVQRDGSGLQL
						LCTVKLQVLPCIIAQGNGGRKA
	Ì					TSSLEAELTSWRPLCLEDGTGF
						RGEDTLETVNAEGRENRRKAG
	İ					HRAAGSSASRRCRNTGRSVQRL
						RPQKTQAFPTHCHFCLILVVKA
						TQLNPKSRVRKVPFAPMKHGK
						ALAGPWANICAGKSSNEIRTCD
	l					RHGCGQYSAQRSQRPHQGVDI
	1					LCSAGSTVYAPFTGMIVGQEKP
						YQNKNAINNGVRISGRGFCVK
						MFYIKPIKYKGPIKKGEKLGTLL
						PLQKVYPGIQSHVHIENCDSSDP
14773	45141	Α	14859	1	2577	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14774	45142	Α	14860	3019	5574	RETRGEVKNSKNQSTSAPPKDC
						SSLADPKKIQTTIREYYKHLYPN
						KLENQEEMDKFLDTYTLPRLN
						QEEVESLNRPITGSEIVAIINSLP
						TKKSPGPDGFTAEFYQRYKEEL
			 -			VPFLLKLFQSIEKEGILPNSFYE
						ASIILIPKLGRDTTKKENFRPISV
						MNIDVKILNKILANRIQQHIKKL
						IHHDQVGFIPGMQGWFNIHKSI
						NVIQHINRTKDKNHMIISIHAEK
						AFDKIQQHFMLKTLNKLGIDGT
						YLKIIRAIYDKPTANIILNGQKL
						EAFPLKIGTRQGCPLSPLLFNIV
	:					LEVLARAIRQEKEIKGIQLGKEV
						VKLSLFADDMIVYLENPIVSAQ
						NLLKLISNFGKVSGYKINVQKS
						QAFLYTNNRQTESQIMSELPFTI
						ASKRIKYLGIQLTRDVKDLFKE
						NYKPLLNEIKEDTNKWKNIPCS
						WVGRINIVKMAVLPKLPMTFFT
						ELEK\TTLKFIWNQKRARIAKSI
						LSQKNKAGGITLPDFKLYYKAT
					İ	ATKTAWYWYQNRDLDQWNRT
						EPSEITPHIYNYLIFDKPDKNKQ
1					ľ	WGKDSLFNKWCWENWLAICR
						KLKLDPFLTPYTKINSRWIKDL
						NIRPKTIKTLEENLGITIQDIGMG
						KDFMSKTPKAMATKAKIDKW
		ŀ				DLIKQESFCTAKETTIRVNRQPT
						KWEKIFATYSSDKGLISRIYSEL
						KQIYKKKTNNPIKKWAKDMNR
	1		1.40.61	100.7	61.40	HFSKEDIYAAKKHMKKCPSSLA
14775	45143	A	14861	1895	6140	
14776	45144	A	14862	1	2875	MKAEIKVFFETNENKDTTYQNL
						WDTFKAVCRGKFIALNAHKRK
						QERSKIDTLTSQLKELEKQEQT
						HSKASRRQEITKIRAELKEIQTQ
	1					KTLQKINESRSWFFERINKIDRS
						LARLIKKKREKNQIDTIKNDKG
						DITTDPTEIQTTIREYYKHLYAN
						KLENLEEMDKFLDTYTLPRLNQ
						EEVESLNRPITGAEIVAIINSLPT
						KKSPGPDGFTAEFYQSWAETQP
		:				KKENFRPISLMNIDAKILNKILA
						KRIQQHIK
L	L	ч	Ļ		L	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14777	45145	A	14863	2843	4460	HQHQKPHPHRNPIQKSPTSKT KVLEVLARAIRQEKQS/GIQIGS QEVKLSLLEDDMIVHLENPIISA KNLLQLISNFTKVSGYKINVQK SQAFPYTNKTQIENQIMSELPFT IATKRIKYLGIKLTRDVKDLFK NNYKPLFKEIREDTNKWKNIPC SWVGRINIVKMAILPKVIYRFN AIPIKLPMTFFTELEETTLKFIW NQKRARIAKSILSQKNKAGGIT LPDFKLYYKATVTKTAWYWY QNRDRDQWNRTEPSEIMPHIYN YLIFDKPEKNKQWGKDS\LFNK WC*ENWLTICKRLKLDPFLTPY TKINSRWIKDLNVRPKTIKTLEE NLGITIQDIGIGKDFMCKTPKA MATKAKIDKWDLIKLKSFCTA KETTIRVNRQPTKWEKIFATYS SDKGLISRIYNELKQIYKKKTN NPIKKWAKDMNRHFSKEDIYA AKKHMKKCSSSLAIREMQIKTT MRYHLTPVRMAIIKKSGNNRC WRRCGEIGTLLHCWWDCKPVQ PLWKSVWRFLRDLELEIPFDPAI
14778	45146	A	14864	1	8335	PLLGIYPKGYKSC MGKKQNRKTGNSKMQSASPPP KERSSSPATDQSWMENDFDEL REEGFRRSNYSELREDIQTKGK EVENFEKNLEECITRITNTEKCL KELMELKTKARELREECRSLRS RCDQLEERVSAMEDEMNEMK QEGKIKRDKEGHYIMVKGSIQQ EELTILNIYAPNTGAPRFIKQVL SDLQRDLDSHTLIMGDFNTSLL TLDRSMRQKVNKDTQELNSSL HQADLIDIYRTLHPKSIEYTFFS APHHTYSKIDHIVGS
14779	45147	A	14865	1	3166	MVKGSIQQEELTILNIYAPNTG APRFIKQVLSDLQRDLDSHTLI MGDFNTPLSTLDRSMRQKVNK DTQELNSALHQADLIDISRTLHP KSTEYTFFSAPHHTYSKIDHIVG SKALLSKCKRTEIITNYLSDHSA IKLELRIKNLTQSRSTTWKLNN LLLNDYWVHNEMKAEIKMFFE TNENKDTTYQNLWDAFKAVCR EKFIALNAYKRKQERSKIDTLTS QLKELEKQEQTHSKASRRQEIT KIRAELKEIETQ
14780	45148	A	14866	1	3912	
14781	45149	В	14867	1	2832	

SEQ ID			SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14782	45150	Α	14868	1944	3531	LRDCKRINQHPVKTDQSAFCK
						M/API/MQDVVLEVLARAIRQEK
						EIKGIQLGKEEVKLSLFADDMI
						VYLENPIVSAQNLLKLISNFSKV
						SGYKINAQKSQAFLYTNNRQTE
						SQIMSELPFTIASKRIKYLGIQLT
		ŀ				RDVKDLFKENYKPLLKEITEDT
						NKWKNIPCSWVGRINIVKMAIL
]					PKVIYRFNAIPIKLPMTFFTELE
						KTTLKFIWNQKRARMAKSILSQ
						KNKAGGITLPDFKLYYKATVT
						KTAWYWYQNRDVDQWNRTEP
1				:		SEITPHIYNYLIFDKPDKNKQW
						GKDSLFNKWGWENWLAICRKL
	ļ					KLDPFLTPYTRINSRWIKDLNV
						RPKTIKTLEENLDITIQDIGMGK
ŀ						DFMSKTPKAMATKAKIDKWDL
1						IKLKSFCIAKETTIRVNRQPTKW
						EKIFATYSPDKGLISRIYNELKQI
						YKKKTNNPIKKWAKDMNRHFS
						KEDIYAAKKHMKRCSSSLAIRE
-						IQIKTTMRYHLTPVRMAIIKKSG
						NN/R/CF*WQKPGPSG*ALRSWK
1						EDIAGTSYVGMNQCQMRKNTK
1						TLI
14783	45151	Α	14869	1182	3891	KMIKGISTTDLTEIQTTIREYYK
						HLYANKLENLEEMDKFLDTYT
		:				LPRLNQEEVESLNRPITGAEIVA
						IINSLPTKRSAGPDGFTAEFYQR
						YKEELVPFLLKLFQSIEKEGILP
						NSFCEASIILIPKPGRDTTKKENF
						RPISLMNIDAKILNKILAKRIQQ
						HIKKLIHHDQVGFIPGMQGWFN
						IRKSINVIQHINRAKDKNHMIISI
						DAEKAFDKIQQHFMLKTLNRL
						GIDGTYFKIIRAIYDKPTANIILN
						GQKL
14784	45152	Α	14870	997	1161	CTSTVIMSICTKLQNKEHVIEAL
						RRAKFKFPGHPK\IHISKK*DFT
						KFNADEFED
14785	45153	A	14871	1	1458	
14786	45154	В	14872	53	1468	
14787	45155	Α	14873	1819	2235	
	I		L		·	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14788	45156	A	14874	707	1613	VRARAGERARVVRENGKAELP
		ŀ				VALVLPSRALSHTFTCTDLKVQ
		ŀ				FHQRLRLQEKGSKFIGSNMSQE
		ŀ				SDNNKRLVALVPMPSDPPFNTR
		ŀ				RAYTSEDEAWKSYLENPLTAA
		ŀ	•			TKAMMSINGDEDSAAA/HRPAL
						*LLQENQSEPIPREGVGWG*GE
		ŀ				ER\ERMTEEEQEGEGGGGRGG
		İ				GEGEEEEEEEDVEEEEEEEEE
						EKEEEEKGGEKEEENEEKEKQR
		ŀ				EGRRGKETSNGQMVRPRLLPP
						HQEHSRNYLVFIMDSNMIEICF
						REASISPKAQAYVLIPDFKNCIY
						GWEEDSIEILWGLPGSHVL
14789	45157	Α	14875	3	346	QRSKSCKVPGLGKPLTLPPKPE
						K\PQGQKDRRPTGFKP*N*RRRR
						SEKSLRFFPPGS/PQAVPIPVGFS
						FDTNL*HFL*KPLIFIMGGTVSL
						PHPRSPSAPPGTPQRPMTFCHLK
		<u> </u>				Q
14790	45158	Α	14876	1868	2666	GLLLSPFQGLQGVAPRKTRSNL
						HPGSLAHLSHQPPAPWPPRGLW
						PPGQETSCHCASGRALGTGSPR
		İ				PPHAPWLHLEPQRAPHSGAPSP
						GSSPSAPLHIPVAPENLQSLGHR
		:				SPPNPSGPDCYPSPSGP/PPSSLD
						SPGLQTAPGAMFLSPRSGVQSA
						CPTPRSLDLRNSPASTPDQSVCP
						TPLSLAWLWPPPSPTSAKSLLFT
						LTSEVQSFSPTPIPPGSSDPPLSK
						QP\SGSCLSSSGHLTSPSVPRHSP
						WPRSSPQNPHSWTLPSS
14791	45159	Α	14877	762	1528	AIPGGAAWPFRAEGAQGPGKR
						LTFKPLGAQRQSVSGGSEVPPL
						PLHLKMAPEMPVAL*SPGTGV
						HPSLLPPT/GPKVPPGRIDRGPVP
					1	TLAPQVGTAPIQWHCRGSRCPS
						G/SRPPGGWR/RPCQAPAPPLGQ
						HLPRLLPPPTGPQGPRPLTPSEP
1						PPWSCWLPASPPAGPGSRRVGG
						GPGTCQGLLRS*GLLLSPFQGL
						QGVAPRKTRSNLHPGSLAHLSH
						QPPAPWPPRGLWPPGQETSCHC
						ASGRAVGTGSRRPRA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14792	45160	Ā	14878	<u> </u>	963	MARKDCCSPPGDSSPGRLPLVL
14//2	45100	^	14070	'	703	PAQCLVLLRLQITAFSSIHVAAN
		ŀ				DMMSFFLMAKQDSIAYKYRIFF
		İ				IHSSVDGHARPLSPSLVVENGC
		ŀ				GKQVFLLCSDNASPEVSTVALA
						SCPGTFPRHCGAAESTHICATG
		ŀ				LGELTPQGHGHIPNKLCACCAT
						GLEELVPRDHGKLGISWFGFIA
		İ				GAYLNLARVVGECQGSYGYLS
		ŀ				LHHYCYCHYYHYCHYHYH\AA
		1				TY*NHYYHYYH/HHNNSYHHH
1						SYYHYSYHFYHYYYNCYHHH
						YHCRHYYHPHDYYHYHA/RHY
						SLYYCHYLCHYYYRCYHDHSS
1						HQHNHHECLP*VRHSAEHLQY
14793	45161	A	14879	105	1448	CHQPPASPFSRREPGGSEKPPER
14775	15101	1	11075	103		GCPAWEPAAQGPVQPLGEDGP
						RLCQPPPPSPDHALG\PYGHQRS
						GRPTLPGPRK\PAKKRGR*PSSS
						TGQ*SHRPTTGTPGGQAGGSG*
1						VTAGGQ*PPQNL*AKISCPEMP
						GPSPGMQQGVKHPSLFSHQTV
						AKGFTTRWNCRQVCPHRGERK
				<u> </u>	,	WISSVRPCGGALAPSCA/DGLR
						GRRTLSGSQPLW/PPAAVSAVG
						QLQPPCHTLTPHPLQNSTQHAE
						PQRGSVYSL*EGTSP\LAAVPFA
						PHIEKRPFSTFPAGDA*PSEAHA/
						PGPACCHSPGGLEGPVSRAGPG
						TSAGR\PGNGRERAWPPHSGPS
						TPPDTSCG/SPGPQLARAGSRRA
						NGSTLPPTRQGFLPGRKDSAGA
				1		CGQP*AEGAAAGPSRSPAAER
			:			AVLPLPR\GRRKCRGGGPGVET
		1				GVSGARRGGDRSGCSPAALDA
						EAPASLAGVSIREG
14794	45162	Α	14880	1	2421	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14795	45162	I .	14881	1080	2571	VHCRFWILALCQMSRLQKSPLL
14/93	45163	Α	14881	1080	23/1	FNIVLEVLAKAIKQEKEIKGIQL
						GKEEVKLSLFADDMIVYLENPI
						V/SAQNLLNLISNFSRVSGYKIN
						VQKSQAFLYTNNRQTESQIMSE
				1		LPFTIASKRIKYLGIQLTRDVKD
				Ì		LFKENYKPLLNEIKEDTKKWK
						NIPCSWVGGRINIMKMAILPKVI
						YRFNAIPIKLPLTFFTDLERTTL
						NFICNQKRAPTLPQVQSLS\QKN
						KAG\GIT\LPDFKLYLQGLQ*PK
		ļ			1	TAWYWY\QNRDIDQWNRTEPS
		1				EIMPHIYNYLIFDKPWEK/HKQ
	:	1				WGKDSLFNKWCWE\NWLAICR
						KLKLDPFLTPYTKINSRWINDL
	1					NVR\PKTIK\TLEENLGIT\IQD\IG
						MGKDFMSNKTPKA\MA\TKAKI
						DKWDLI\KLKSF\CTAKETTIRV
						NR\QPTKWKKIFTTYSSDKGLIS
		ļ				RIYNELKQIYKKKTNNPIKKWA
						KDMNRHFSKEDIYAAKKHMK
					:	KCST\SLAIREMQIKTTMRYHLT
		ŀ				PVRMAIIKKSENSRCWR
14796	45164	Α	14882	136	420	RESRNLSRGEESEDPAEPS\RNG
						*EHPLCYLQ*EFLQVLTMLQAE
		ŀ				GTMHHFRSICQVNRNFLERGH/
						SPPSPAPPPETHTGSPRPPSGRSR
		ŀ				IRAYLH
14797	45165	В	14883	1	1197	
14798	45166	A	14884	3	538	GAGTDGGRTPAPRAFPDPLGEA
				-	-	RTSAPTGPARPGGSASAARPAA
						VRPAEPAAEETQEQPRAGKRAI
						HEYRYVYVQQRRKTDPQPAVA
						VFGIPADGD*VSRTAYHAHQPL
						AWSGR/E*MCLSMAMWNT*ST
1						MKK*ISIRVISRCSGPVHRTN*QI
		1				PEPVRAWRFLICRCICFSPGRWI
						KT
ł	L	1				K1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14799	45167	Α	14885	1208	2765	KMAPTWDKMVSSAQMGFNLQ
						ALLEQLSQDELSKFKYLITTFSL
				:		AHELQKIPHKEVDKADGKQLV
						EILTTHCDSYWVEMASLQVFEK
						MHRMDLSERAKDEVREAALKS
						FNKRKPLSLGITRKERPPLDVDE
						MLERFKTEAQAFTETKGNVICL
						GKEVFKGKKPDKDNRCRYILK
						TKFREMWKSWPGDSKEV/RGY
		1				G*EIQDADPIQQPQGASRAL\PY
						TVVLYGPAGLGKTTLAQKLML
ŀ						DWAEDNLIHKFKYAFYLSCREL
						SRLGPCSFAELVFRDWPELQDD
						IPHILAQARKILFVIDGFDELGA
						APGALIEDICGDWEKKKPVPVL
						LGSLLNRVPLCIENKLGVGGED
				[WGDQLRVYWWNPSKREGGSG
		1				WSHGSEDANQSLKHLRLSANV
1						LLDEGAMLLYKTMTRPKHFLQ
		ļ		<u> </u>		MLSPWETTLAQIIMLDWAEETS
		l				SDKLKYAFYCSCRELSRLCPCN
						FEELVFRDWPELQDDIPHILAQ
						ARKILFVIDGFDPLGGRTLWTL
						TRGHLRELGNRRKRCPSSWGV
14800	45168	Α	14886	213	625	EPGIDTSCLCAGRCSRPGKGRA
		ŀ				CGGSSPCSWPGQPPAPPAAAAV
		ļ				PACPSGGRGAGPIQSGTCCSRS
						ADSRCTSWTAW/WPPGLGAAT
ł						VQTVMS*TV/WPMPHVPQAGW
]				LALGQGAAAAKHPLTSATTRS
		ł				WCLGVGPGPWM
14801	45169	Α	14887	1	375	MCNAQVLEYMGKSSSLLTSDL
		l				QLVRDALRSLRNSFSGHDPQHH
		ŀ				TIDSLEQGISSLMERLHVMETQ
		ŀ				KKQERKPPLVIPRQT/WVWSGP
						PANSNRPAAEGPDC*KEN*RDA
						LRSLRNSFSGHDPQHHTIDSLEQ
						GISSLMERLHVMETQKKQERKP
]		PLVIPRQTGSGVDLQQTPTDLQ
						LRDLTVRRKTNEQKGIASASTK
						RTSTPKPHL

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NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ł	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14802	45170	Α	14888	1	750	MGELNQGSWRQGHLPLDAAN
j						GIAIRVPRRRKDESRKNPTRKPH
1						SWKPTKNFEDDVEVMKRSMSI
1						RIEDTHSGPSAAGVLEFAGGPL
						QTLFAWVSPAEAAEQQRLLPVP
						/CLWKLCPRGAPARC\RQNSPSK
						LPSGFVYTVRGKLPTQASVVVD
						APPATKLECPRSTSDCCAGSKN
						FKPVDLSLLASVGPRVNLQQES
						SHYGKISVTVAGHCPTHATGRI
						LGKQALRWSPMCSMFIGRALG
						LVPVEGGLLQIA
14803	45171	Α	14889	81	164	VGKQGLEWTPANSSRPAAEGP
			. =			DC*KEN
14804	45172	Α	14890	1	674	MVVYACSPSYLEAEVGGSLSLE
ŀ			ŀ			VEATNPKNSSKKFLDLISEFCK
ļ						VSGYKISVHKSVALLYTKNDQ
						AATQIKNLIPFTTAAKQFKYFGI
						YLTKEEERLKPSRKKITKKHTK
						KRTASLILHAMICSQIPKQQQNE
						KYQVPQFDQSTIKNIESAKGLD
						VWDSWPLQNADGTPPLVIPRQ
						T/WVWSGPPANSNRPAAEGPDC
						QKEN*QTERNSINIKKKDIHTKT PLCI
14805	45173	A	14891	619	931	YLLMDFPLLHSWLVSLGDKVP
1.000					1.0	HEFWRGCPSCSFGPRFWVAGV
						LVL*APKL*LRQVRSAGKRCGQ
						KW*IK*/SDRLCVD*NK/WSDSD
	İ					LKPATSCSLIVTCHSLLGF
14806	45174	В	14892	1	658	
14807	45175	Α	14893	48	288	LCSIKKPEKSLLCW*YPGNRVW
		ĺ				SGPPAKSNRPAAEGPDC*KEN*
						QTERNSININKKDIHTKTPSEGH
						QHQRPKTKPGRI
14808	45176	A	14894	801	1157	QSGPSAAGLLDFAGGPLQTLFA
						RVSPAEAAEQQRLLPFPSSGKL
						RPRGAPARCQPELSCMRC\R*SL
						LGGVSQSGYMGVRDPFEEAAC
						PLSELECSAGRSAALLRMVRQG
			<u> </u>			HLGLLKLRP

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NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14809	45177	A	14895	1	1891	MPVWVTVVKSWNTQIGRPGH
ŀ						VSALGFELTKNPWPLTYESGDG
						RSLGKPGAVSRRGGRHGPHPQ
						QAECDLPAGRIAGPCERGLERQ
						EKLPPAPQGGAWHLRLSRERFK
						SEDAPKIHVALGGSLFLLNLAF
						LVNVGSGSKGSDAACWARGA
						VFHYFLLCAFTWMGLEAFHLY
						LLAVRVFNTYFGHYFLKLSLVG
						WGLPALMVIGTGSANSYGLYTI
i						RDRENRTSLELCWFREGTTMY
						ALYITVHGYFLITFLFGMVVLA
						LVVWKIFTLSRATAVKERGKN
						RKKVLTLLGLSSLVGVTWGLAI
						FTPLGLSTVYIFALFNSLQAHKQ
						LKDRKVGPHFMQGAVFLEGKF
İ	<u> </u>					WGVDIASFEPQEQFWWTNLGK
						EDEDKSFDMPHSWVEQIEISPEE
ľ						EAMGSAQMVRGVSGPERGSKK
						RPQLTVSTAFETRCPNGKKVIQ
						YKRAKLEKWAPYLNSNGLVSR
						LTTYEDLQCTNILEIKEWYQNR
						EDMLELKHINKTTDLKTDYFKP
						GHPQALRVHSYKSMQPEMDRV
						IEFYETARVDGLMKREETPRTM
						TEYYQGRPDFLSYRHASFG/RPS
1						QEAHSEQCRVKPPAHCAKPAE
						DDV/RKSACFWSRRSASSCATT
						AVRTTSRPPSASSCGAPRWTAK
14810	45178	Α	14897	1	425	HVQQTEISRTVSTGSPIILSCSVP
						VTLPNGPVLWFKGN/AGPNRKL
						IYNFKQGNFPRVKEIGDTTKPG
						NTDFSTRIREISLADAGTYYCV
						KFIKGRAIKEYQSGRGTQVFVT
:						VSYSSRSQKSEICISGLKSRHQQ
						NCSFWRF
14811	45179	Α	14898	J	393	RQGNFPRAKQIGDTTKPGNTDF
						STRIREISLADVGTYYCVKFIKG
						RAIKEYQ\SGRGTQVFVTEQNP
						RPPKNRPAGRAGSRAHHDAHT
						CLSALPERNSTNYFVQPCCCLR
						LLGLTGFAVKIIQTGKERTSK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14812	45180	A	14899	1	753	MGSSLSPVLYVALPVRQPLLRS
14012	75100		14077	1	755	VVEERSTKMRVGSPGFQAKFRF
						YSQGLSWSARGEEGLVPAFWL
						QPYPRPLNFKAEMTEQKENQT
						YVDGGFRPYIHRRKNLTEGHK
			İ			AEKEMEPWDGFSDESCRTHCQ
						VHDYAKGHGEVQRERVDEQK
						EHLGGRRQDCQQQHLNRRLHC
						/TVSGEAPSEASGSVRGHGRAH
						GLAGSERHGHDHAAPEHQLLQ
						SVDADHVVPVAWYQFWDDEG
						RGILSPGDHKPDRRGAALY
14813	45181	A	14900	6	726	VEGISRTGIQAEQAPGFTTMPIP
14613	43161	A	14900	0	1/20	VSVNPGDVLTLACNMSALSPE
		l				GPVLWFKSIGPGQQLIFSCNGS
						HFPRVTPVENTMVDQTDYSIILS
		İ				CSVPNTLPNGPVLWFKGTGPNR
		ĺ				KLIYNFKQGNFPRVKEIGDTTK
						PGNTDFSTRIREISLADAGTYYC
						VKFIKGRAIKEYQSGRGTQVFV
						TGEYVSYTPYPLVYNS\PLWDE
			İ			ASRGRSRQQSLLFCSLHW*YPR KQDLEWTSSKLQQTCRRVA
14814	45182	A	14901	760	978	KKLLLLLCGIEWHGFSSDSGS
14614	43102	l'A	14701	1,00	978	GGWVPEG*K*GCEHELLPEDVP
		1				SCFGWAECWLVRRTLPLFHYV
		1				ELCYSFHG
14815	45183	В	14902	772	1165	ELETSITIO
14816	45184	A	14903	1	2133	
14817	45185	A	14904	11	1311	MATTPGRALGSLLPTLLLGLTA
						GQPFISLTGPSHRTSPGDSVPFN
						CTAVPFNSQDFSRDFNVTWLK
						DSDEHPASAQRLVPDNGGNDFI
						TSKAWVTLTRODVSSEITCEVT
						HRALAEPLKTTMNLSQATRNP
:			ŀ			DGTYSLEHTWQAEATLEEREF
						ACWVVHDEQPPLKANVTLQAQ
						VRRQGKGQSVQPYKLQGPLQR
						SEPGTRIRLTYASSGFSTHQVTV
		1				TWLKNKHELPNPQTSVQYSGH
		1				TYNVTSSVLVPLMDDDVFSHV
		1				VCHVEHKLTWFFQKTTGLDQY
		1				L/QG/DP*QLCGGHLTLPSSPRAP
						PQLPGLQHP/VVFHSQVLLFSCL
						VAVTCLVQRFYPQNVHLAWLE
						DCHTLKGTEQPTLKKNNDRSY
		l				TLEKLLLVNASVQGPERVLTC
						MMEHEGQPPKRANLVLSTAAH
			1			TAYKPIGSTESIMVTMIITISQM
L	<u> </u>	Ь.	L	I	<u> </u>	TATRIGSTESIM V TWITTISQIVI

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14818	45186	A	14905	3	341	RRATRPGN*GGRHNAGRSSARL ETGRPAMAPQGRPRPLGCRTPQ HPGHWPSAPGLLPPAPGPVAG MTQAQRWCPEPDAHFLQYTLY FVRHIPQLPKQTPHHYIATRLPC ATQ
14819	45187	A	14906	154	679	GVCCPLLGGVSS*ATWGSGTT* GGNTHLI
14820	45188	В	14907	1	2554	
14821	45189	A	14908	387	826	ARAACLPRGSARGTAGAGEEPP DPAPLHLRGEGPAP/HRPRSPRG MGWPAAPSTRSPPST*RRCPLPP RPRWSP*SHPGPRASARRLP*LS *RASRSSEAPSPGSLGSVAGQD CSRTGTGFSPPAFPSDGQQTPPP APSSVPTLNSLG
14822	45190	A	14909	86	2223	GQEPQVREQKACHHLGSSPPPS WEL*EQRPAGP*N**PSTLTTGT GSTTT*QIRMKVN*MGN/DSPQ GKNTPKMYSGEFSPVRVHVPFS LSDLKQIKIDLGKFSDDPDGYID VLQGEDILAKAGAIRHLNIGEG THVCCPLLEEGINPEVWATEGK YGRAKNAHPIQVKLKDSASFPY QRQYPLRPEAQQGLQKIVKDL KAQGLVKPCSSPCNTPILGVQK PNGQWRLVQDLRIINEAIVPLY PAVPNPYTPLSQILEEAEWFTVL DLRDAFFCIPVHPDSQFLFAFED PLNPMSQLTWTVLPQGFRDSPY LFGQALAQDLSQFSYLDTHVL WYMDDLLLAARSETLCHQATQ ALLNFLATCGYKVSKPKAQLC LQQVKHLRLKLSKGTRALSEER IQPISAYPHPKTLKQLRMFLGIT GFCQIWINRYGKIARPLYTLIKE TQKANTHLVRRTSLQPVAYLS KETGTSLQPVAYLSKEIDVVAK RWPHCLRVVAAIAVLVSKAVK MIQGRALTVWTSHDVNSILTAK GDLWLSDNHNEEKIEHNWQQV IAQTYAAQGDLLEVPLTDPDLN LYTDGSSFVEKGLRKAAYAVFS DNGILESNPLTPGTSTQLAELIA VTQALELEGKREPENPADNASY SCEPLEDLRLLFRRQPIEAVKLQ MVLQMEPQMQSMTKIYRGPLD WPASPCSNVDDIEGTPPKEISTA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14823	45191	A	14910	929	1324	RRYQVQRPSEGTLVYLFWPV*/ PVVGVDPWLPTGPD\YSLLVPIP PPRARTPRWA*A*MKGCRRSAT A*WCMC*SCRVPCPSCTPMFHS PCSAMTAVTPSLARP*THGSPP KAQGAPQGVKGPSSGLEAPPW A
14824	45192	Α	14911	2	286	
14825	45193	Α	14912	220	249	P*N**PSTLTTGTGSTTT*QIRM KVN*MGN/DSPQGKNTPKMYS GEFSPVRVHVPFSLSDLKQIKID LGKFSDDPDGYIDVLQ
14826	45194	A	14913		163	EPENPADNASYSCEPLEDLRLL FRRQPIEAVKLQMVLQMEPQM QSMTKIYRGPLDWPASPCS*DF LGRGAFNVINIGAWASRPVQGS AVDLSHGLHLGLHLKNHL
14827	45195	Α	14914	1	287	
14828	45196	Α	14915	2498	2788	
14829	45197	Α	14916	112	333	
14830	45198	Α	14917	329	1666	
14831	45199	Α	14918	1	756	
14832	45200	Α	14919	1256	1651	RRYQVQRPSEGTLVYLFWPV*/ PVVGVDPWLPTGPD\YSLLVPIP PPRARTPRWA*A*MKGCRRSAT A*WCMC*SCRVPCPSCTPMFHS PCSAMTAVTPSLARP*THGSPP KAQGAPQGVKGPSSGLEAPPW A
14833	45201	A	14920		987	MLVLGYNRKNTEGTQKQKGT NASDFHFLSQVLEQVVSPKGSK EAQCCVLRHLGCESSESAPGIPP NLGIQLLTWAVMWDPFPTTLA RAPSLALELMTQYFNNWNWV YNNITDQGESKMSKLKGKERR QREGERRERKREREKRERESQR KERKREKKGKKREREKDRSDL KQIKIDLGKFSDTPDGYIDVLQ GLGQPYYLTWRDIMLLLDQTL TPNERSAAITAVREFGDLWYLS QVNDRKITEEREQFPTGQQAVP SVDPHWDTESEHGDWCHRYLL TDVLEGLRKTRKKP\IN*SMISTI TQGKEENPTAFLERLREALRKH
14834	45202	Α	14921	473	1015	
14835	45203	Α	14922	3	937	
14836	45204	Α	14923	530	612	LQAKVFFLLRWPAILRG*NYGP RGVPH
14837	45205	Α	14924	101	430	

AQPARSAAKPGLDQPGSHARL RGRGGEARQSLEAIRGGPSGIA PSDIPLPAKPSRRVIAQTYAARG NLLEVPLTDPDLNLYIDGSSFVE KRLRKAGIKAVKLQIVLQMEPQ MQPMTKIYCGPLDQPASPRSNV DDIEGTNASDPFLSQVLEQVV SPKGSKEAQCCVLRPLGCESNL KQIKIDLGKFSDTPBGYIDVLQ GLGQPYYLTWRDIMLLLDQTL TPNERSAAITAVREFGDLWYLS QVNDRKITEEREQFPTGQQAVP SVDPHWDTESEHGDWCHRYLL TDVLEGLARTRKKPJIN*SMISTI TQGKEENPTAFLERLEALRKH TQKEENPTAFLEALRA TQKEENPTAFLEALRA TQKEENPTAFLEALRA TQKEENPTAFLEALRA TQKEENPTAFLEALRA TQKEENPTAFL	SEQ ID	SEQ ID NO:		SEQ ID NO:	2		Amino acid sequence (X=Unknown,
14838	NO:	1	hod	1			
14838 45206		sequence		09/540,217		or peptide sequence	detetion, \=possible nucleotide insertion)
PRRHRYKKKWAATEPKFPAVR					sequence		
LALQNFDMTYSVQFGDLWPSIR VSLLSEQKYGALVNNFAAWDH PGDIVLDLCAAPGGKTLALLQT GCCLTWSRKRCAALSMFSRPA RCLCRARTTARRAVRRPDQRR ATAAATKQQRARTAPEAPGSSA HSSYCTMIPSACCLPRQLKPH AQPARSAAKPGLDQPGSHARL RGRGGEARQSLEAHRGPPSGIA PSDIPLPAKPSRRVIAQTYAARG NLLEVPLTDPDLNLYIDGSSFVE KRLRRAGIKAVKLQIVLQMEPQ MQPMTKIYGGPLDQPASPRSN DDIEGTNASDFPFLSQVLEQVV SPKGSKEAQCCVLRPLGCESNL KQIKIDLGKFSDTPDGYIDVLC GLGQPYVLTWRDIMLLDQTL TPNERSAAITAVREFGDLWYLS QVNDRKITEEREQFPTGQAVL SVDPHWDTESSHGDWCHRYLL TDVLEGLRKTRKKPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKKPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKKPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLERLREALRKH LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLEGLBARAT LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLEGLBARAT LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLEGLBARAT LTDVLEGLRKTRKCPIN*SMISTI STENSLATARAT STENSLATARAT STENSLATARAT LTDVLEGLRKTRKCPIN*SMISTI TQGKEENPTAFLORG LTDVLAGLRHGHTTI TTPNLTATARATARATARATARATARATARATARATARATARAT	14838	45206	Α	14925	1	1587	MAALTLRGVRELLKRVDLATV
VSLLSEQKYGALVNNFAAWDH							PRRHRYKKKWAATEPKFPAVR
VSLLSEQKYGALVNNFAAWDH							LALONFDMTYSVOFGDLWPSIR
PGDIVLDLCAAPGGKTLALLQT GCCLTWSRRRCALASMFSPR RCLCRARTTAPRAVRPPOQRR ATAAATKQQRATPEAPGSAA HSSYCTMIPSACCLIPRQLKPP AQPARSAAKPGLOQPGSHARL RGRGGEARQSLEAIRGGPSGIA PSDIPLPARPSRRVIAQTYAARG NLLEVPLTDPDLNLYIDGSSFVE KRLKRAGIKAVKLQIVLQMEPQ MQPMTKIYCGPLDQPASPRSNV DDIEGTNASDFPFLSQVLEQVV SPKGSKEAQCCVLRPLGCESNL KQIKIDLGKFSDTPDGYIDVLQ GLQQPYYLTWRDIMLLLDQTL TPNERSAAITAVREFGLWYLS QVNDRKITEEREOFPTGQQAVP SVDPHWDTESEHGDWCHRYLL TDVLEGLRKTRKKPIN*SMISTI TQKEENPTAFLERLEALRKH TQKEENPTAFLERLEALRKH TQKEENPTAFLERLEALRKH TQKEENPTAFLERLEALRKH TQKEENPTAFLERLEALRKH TQKEENPTAGGETSSSSGOLDRKKPLFSL EFGSPGETEDKSRQRQDAGSPK SEDTPAGGFTNSSSSGODSDKTK FPFSLGLGAPGKAEDKSGOSQD AGGSKSEDTPPGGFFYSSSSS SSDKKKPLFSFEFGATGEDEDK SRERWDAGNSRSEDS TABABA 45209 B 14928 110 269 TABABA TABABA TABATT TOKATA TOK							-
RCLCRARTTAPRAVRRPDQRR ATAAATKQQARATPEAPGSAA HSSYCTMIPSACCILPRQLKPPH AQPARSAAKPGLDQPGSHARL RGRGGEARQSLEAIRGGPSGIA PSDIPLPAKPSRRVIAQTYAARG NLLEVPLTPPDLNLYIDGSSFVE KRLRKAGIKAVKLQIVLQMEPQ MQPMTKIYCGPLDQPASPRSNV DDIEGTNASDFPLSQVLEQVV SPKGSKEAQCCVLRPLGCESNL KQIKIDLGKFSDTPDGYIDVLQ GLGQPYVLTWRDIMLLLDQTL TPNERSAAITAVREFGDLWYLS QVNDRKITEEREQFPTGQAVP SVDPHWDTESEHGDWCHRYLLL TDVLEGLRKTRKKPIN*SMISTI TQGKEENPTAFLERLREALRKH 14839 45207 B 14926 I 939 14840 45208 A 14927 I 4317 MQSMTKMYCGPLDQPASPCSN VDDIEVKLEIMLLTERDDGKEV LDVGEERRYEIEEYDSKWTRK YNCLAVLVSHSSQQITTMKLIL WYLVVALWCFFKDGKIAASRS GGFSYGSSSGDLDRKKPLFSL EFGSPGETEDKSRQRQDAGSPK SEDTPAGGFFNSSSSGDSDTK PFFSLGLGAPGKAEDKSGDSQD AGGSKSEDTPPGGFFYGSSSSG DSDKKRPLFSFFGATGEDED SRERWDAGNSRSEDS 14841 45209 B 14928 I10 269 14842 45210 A 14929 I 561 EVFPACSQLVYWGKPITIYPLCE NNVYMLSPNASVCLYSPLAEQ FSHQFPSHDLPSVLAKFSLVSL SEFERNPLAPAVQETQLIQMVV WMLQRRLIQLHTYVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT							1
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SEDTPAGGFFNSSSSGDSDRTK PFFSLGLGAPGKAEDKSGDSQD AGGSKSEDTPPGGFFYGSSSSG DSDKKKPLFSFEFGATGEDEDK SRERWDAGNSRSEDS 14841 45209 B 14928 110 269 14842 45210 A 14929 I 561 EVFPACSQLVYWGKPITIYPLCE NNVYMLSPNASVCLYSP\LAEQ FSHQFPSHDLPSVLAKFSLPVSL SEFRNPLAPAVQETQLIQMVV WMLQRRLLIQLHT\YVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT							GGFSYGSSSSGDLDRKKPLFSL
PFFSLGLGAPGKAEDKSGDSQD AGGSKSEDTPPGGFFYGSSSSG DSDKKKPLFSFEFGATGEDEDK SRERWDAGNSRSEDS 14841 45209 B 14928 110 269 14842 45210 A 14929 1 561 EVFPACSQLVYWGKPITIYPLCE NNVYMLSPNASVCLYSP\LAEQ FSHQFPSHDLPSVLAKFSLPVSL SEFRNPLAPAVQETQLIQMVV WMLQRRLLIQLHT\YVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT							EFGSPGETEDKSRQRQDAGSPK
AGGSKSEDTPPGGFFYGSSSSG DSDKKKPLFSFEFGATGEDEDK SRERWDAGNSRSEDS 14841 45209 B 14928 110 269 14842 45210 A 14929 1 561 EVFPACSQLVYWGKPITIYPLCE NNVYMLSPNASVCLYSP\LAEQ FSHQFPSHDLPSVLAKFSLPVSL SEFRNPLAPAVQETQLIQMVV WMLQRRLLIQLHT\YVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT							SEDTPAGGFFNSSSSSGDSDRTK
DSDKKKPLFSFEFGATGEDEDK SRERWDAGNSRSEDS 14841 45209 B 14928 110 269 14842 45210 A 14929 1 561 EVFPACSQLVYWGKPITIYPLCE NNVYMLSPNASVCLYSP\LAEQ FSHQFPSHDLPSVLAKFSLPVSL SEFRNPLAPAVQETQLIQMVV WMLQRRLLIQLHT\YVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT							PFFSLGLGAPGKAEDKSGDSQD
SRERWDAGNSRSEDS 14841 45209 B 14928 110 269 14842 45210 A 14929 I 561 EVFPACSQLVYWGKPITIYPLCE NNVYMLSPNASVCLYSP\LAEQ FSHQFPSHDLPSVLAKFSLPVSL SEFRNPLAPAVQETQLIQMVV WMLQRRLLIQLHT\YVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT							AGGSKSEDTPPGGFFYGSSSSG
14841 45209 B 14928 110 269 14842 45210 A 14929 1 561 EVFPACSQLVYWGKPITIYPLCE NNVYMLSPNASVCLYSP\LAEQ FSHQFPSHDLPSVLAKFSLPVSL SEFRNPLAPAVQETQLIQMVV WMLQRRLLIQLHT\YVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT							DSDKKKPLFSFEFGATGEDEDK
14842 45210 A 14929 1 561 EVFPACSQLVYWGKPITIYPLCE NNVYMLSPNASVCLYSP\LAEQ FSHQFPSHDLPSVLAKFSLPVSL SEFRNPLAPAVQETQLIQMVV WMLQRRLLIQLHT\YVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT							SRERWDAGNSRSEDS
NNVYMLSPNASVCLYSP\LAEQ FSHQFPSHDLPSVLAKFSLPVSL SEFRNPLAPAVQETQLIQMVV WMLQRRLLIQLHT\YVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT	14841	45209	В	14928	110	269	
FSHQFPSHDLPSVLAKFSLPVSL SEFRNPLAPAVQETQLIQMVV WMLQRRLLIQLHT\YVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT	14842	45210	Α	14929	1	561	EVFPACSQLVYWGKPITIYPLCE
SEFRNPLAPAVQETQLIQMVV WMLQRRLLIQLHT\YVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT							NNVYMLSPNASVCLYSP\LAEQ
WMLQRRLLIQLHT\YVCLMASP SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT							`
SEEEPRPREDDVPFTARVGGRS LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT			ŀ				SEFRNPLAPAVQETQLIQMVV
LSTPNALSFGSPTSSDDMTLTSP SMDNSSAELLPSGDSPLNQRMT							WMLQRRLLIQLHT\YVCLMASP
SMDNSSAELLPSGDSPLNQRMT							SEEEPRPREDDVPFTARVGGRS
							LSTPNALSFGSPTSSDDMTLTSP
I I I I I I I I I I I I I I I I I I I							SMDNSSAELLPSGDSPLNQRMT
LINDEASESEIT	L		L				ENLLASLSEH

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14843	45211	A	14930	1891	2522	GLSNQNQQPGRCIPDAHPVQDL VQDLVHTWCRVPCTSYVPCV* VPGAYLVHTRCIPGAGPGAYPV HTWCRTWCIPGAGFPAL/RPYH ALLLLSDEKSLLGELPIDCSPAL VRVIKTTSAVKNLQQLAQDAD LALLQAMGSAVA\LVMWAWP RATGPLKYCLRVGAGGACGWR PRLVAVSRALLLVWRCGSPWG PALAWQKSPEYSTKVLEKQ
14844	45212	С	14931	113	274	TALAWQKSIETSTRVLERQ
14845	45213	A	14932		1634	MEIPAVILTVAPILHLTSCGFPV TGDGLKQAEGKCAPEEWNPTK FIKLTDNQACFACGLWPLSEIQ RRTSRTVTIISRISALREKGNGR KPTSISGSVNEMRQVLVKDQVS MPRVLPTIIFVPDEAGFSAGAV WPCAAPLWGSSWTPLKVENCS FYIRLSFAISVGGDELKIAKLVS AFPERAYDLVGHQHQRPKVDK TTKMGRNQSRKAENSKNQNMS SPPKEHNSSPAREQNWMENEF DELREVGFRRSVITNFSELKEH VLTHHKEAKNLEKRLDEWLTR INSVEKSLNDLMELKTTLRELR EAYISFNSQFDQAEERVSVIEDQ ISEIKQEVKFREKRVKRNEQSL QEIWDYVKRP\NLRLIGIPESDE ENGTKLENTLQDIIQENFPNIAR QANIQI/QGKLQRTPTKVASSRT ATPRHIISVGSPKLNMK\EKILR AAQRERSGLPTKGKPFRLTAAD LSAETSTSQKTEWGPIFNILKEK NF\QPR\ISYPAK\LSFISEGEIKYF TDQAKCLRDF\VTTRAFSCLP*K STRPALKELLKGST

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
į				sequence		
14846	45214	I A	14933	1	1429	MVLAAGDPVEKGITWARENAG
14040	73217	`	14733	1	1727	IYISCTMMKLSMIPIGDQSYRNL
						IPLKFVACILIEGNAKCWPLPEF
			·			DPSQIRLIVYQDCERRGRNVLF
						1
						DSSVKRRNEDISVSRIKYLGIQL
						TRDVKDLFKENYKPLLNEIKED
						TNKWKNIPCSWIERINIVKIAILP
						KKLGSDAQVKVFGKCCQLKPG
		1				GDSSSSLDSSVTSSSDIKDQCLK
						YQRTVDLSAQLSSSGKGQTTSS
						SGSLTPMTPHWEIPPSGGQQTS
		1				HTGELRLASGRGSSGMKLPEEG
		1				TDRNLCCSAASSGDTQANRDW
						SGPPANTSRPAAEGPDCDRENG
•						TKLENTLKDIIHENFPNIARQAN
						IKIQEIQRTQQRYCSRRATPRHII
					1	VRFTKVEMKEKMLRAAGEKH
						WVIHKGKSIRLTVDLSAETLQA
						RREWG\PIFNVLKEKNFQLRISY
l					-	PAKLSFISEGEMKSFPEKQMLR
						DFVTIRLALQELLKEALNMARE
						/NR*QPLQKHM
14847	45215	Α	14934	2320	2811	ESKLTRGVKDLFRENYKPLLNE
						IKRGHKQMEEHSMLMGRKNQ
ļ						YREKWPYLPKVNL*IQCHPHQA
						TNDCFHRIGKKLL*SSYGTKKE
		1				PALPSQS*DKRTKLEASHYLTS
						NYTTRLQ*PKQHRYWYQNRDI
						DOWNRTEASEIMPHIYTILILWT
						NLRKTSMGERHP
14848	45216	A	14935	2	872	
14849	45217	Α	14936	769	828	
14850	45218	В	14937	1	669	
14851	45219	Α	14938	1	807	
14852	45220	В	14939	35	212	
14853	45221	Α	14940	235	837	QQAHKTPQRRHSAFTEYVRNN
	1	1				RLPETVIRVKQPALAIFSPDIAPL
						FVHFRADDDVTARVIHAIIMIF
						WCDRVEKRVSDAVACFTLESEI
						AEVGVLCRTHIRQ*RQVTPARM
						*ALSRILHACRSEDVRHNRPPYP
						VIRVKQPA\RRDLAPT*PHCSSIS
						AQTITSLPGCMREVT/ELRPEFF
						K*RKTVLRPTPIMRARAERRCK
14854	45222	С	14941	1	1755	
14855	45223	A	14942	1	747	
14856	45224	В	14943	1	2466	
			<u> </u>			

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14857	45225	A	14944	1	333	LNLVLFQQRAELMIS*PQ*FSRA LLVIFPLLQTELQQLFFLLHVNP AAVEAPPSVPIGLYSENVRNNR LPETVIRVKQPALARFSPDIAPL FVHFRADDDVTARRVEKRCK
14858	45226	В	14945	1	1794	
14859	45227	A	14946	1183	1899	LNRRDSYVEYLADKVADREED AEEYLEAIMEARVTVAGMGLV MEVQDYFDGEADRLAKAWLA EYTPQIKSLKDERKEAYRQIVE MSTEPQDVDLVRPANKFEMTR VREGEKEADLPVWKHHL/SV*R KRELSGSVEPLGNQVFEIETKRE NS*EL**KSLKDERKEAYRQIVE MSTEPQDVDLVRPANKFEMTR VREGEKEADLPVWKHHLLCDE SGNYPALLNHWETRFLRSKPN VKIHRSSDKSPTYFVFFISQYDL PAFYPINGHFLVERVVKVRIVFI EEFADHHHLAIFQFNNRLQRHA FVVTLANVTNQPFSVVVVVNL
14860	45228	В	14947	1	3099	
14861	45229	Α	14948	1174	1545	
14862	45230	Α	14949	1	945	
14863	45231	Α	14950	62	576	
14864	45232	A	14951		1642	MLANIGFYNSNVIHRSKGAIGE GPPDPRGIVEQNLGGDISIPCYN GPRPRRSAQKTTLGPLNQPRIPS AQCAGAFPRVAAAGAGGGG GGSGGSGFCLGWRLRRRRQRR RRWRRGEGEGDGDAGHMRPR PAGRAEARRRRLRGNPPPGAR LPNREGNGSGVSWAAEVSGDG RGARTEAARTRRPGGADNYP GAVPPVQLLTGWCVTAKAPPDI SAISALTAVKHGNCSSLTPLLNP PGSDVIVCAEMDEQWGYVGAK SRQRWLFYAYDSLRKTVVAHV FGERTMATLGRLMSLLSPFDVV IWMTDGWPLYESRLKGKLHVI SKRYTQRIERHNLNLRQHLARL GRKSLSFSKSVELHDKEELAAL FSELKQEQKKVDELIAKLVKNR TRIVNESDVFSWVIRREFQELR HPVDEEKARCLEGIGGHTRGLV ASLDMQLEQAQGTRERLAQAEN LCWNSSEMRTTMSSSGSSTPWP PVSRRPWTERRLGSVRGDRLR ERSRPAVNRTPRMSRVPSGAPR GTPSRRWWMMKCWDHQMRG LVEEEVG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14865	45233	Α	14952	11	1513	MIIALQKESLMSLQAQLDKALQ
						KEKHYLQTTITKEAYDALSRKS
						AACQDDLTQALEKRTGRDWNP
						PPFRTQQRGSKSLTAIRPLCLFR
						KLNHVTSETKSLQQSLTQTQEK
						KAQLEEEIIAYEERMKKLNTEL
						RKLRGFHQESELEVILYEEEMG
						NHNENTGEKLHLAQEQLALAG
						DKITSLERSLNLYRDKYQSSLS
						NIELLECQVKMLQGELGGIMG
						QLSKTSSDLSKM/RCY/TTW*M
						WTKLQFIHSILTSVIFHLLSFSSM
		ł				GSI*KWIMALSGSSFLLDAHYIS
				1		QFGPRQLPSVCAEPEKEKATDG
				1		VVRNGKSTAGHQRYLCSHCRK
						TWQLQFTYTASQPAGQ*RHRL
						RG\MDEQWGYVGAKSRQRWLF
						YAYDSLRKTVVAHHREQGSIK
						CKLEEDLQEATKLLEDKREQLK
	i	İ				KSKEHEKLMEGELEALRQEFK
[ł				KKDKTLKENSRKLEEENENLR
			:			AELQCCSTQLESSLNKYNTSQQ
		ł				VIQDLNKSPMEIRFQLYIFTPDQ
14866	45234	Α	14953	125	363	INVSVWSQPFPGG*QPEKGLKE
						HPADARPGSC\PPAGRPVQGRA
						EVASAAPPRGSSPAAAPGQSHP
						PSCLRAPLYWSFSR
14867	45235	Α	14954	306	583	TCSTRRKNDLSVKYVSVWSQ
l		1				PFPGG*QPEKGLKEHPADARPG
		Ì				SC\PPAGRPVQGRAEVASAAPP
		ŀ				RGSSPAAAPGQSHPPSCLRAPL
						YWSFSS
14868	45236	A	14955	225	407	LRGMVTFSPFSSILNGQFEPNSG
						CPL/CPVCWVCLREPGDPEKLG
						EFLQKDNISVHYFCLVS
14869	45237	A	14956	484	722	NTCRCGHNRSPEVNSRRGGLRS
						IPRMPDAGSCS/PPAGRPSAGPG
						RRSPPPLLPRGSSPAAAPGQSHP
						PSSSQGAAVLEL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		07/340,217	sequence	or pepade sequence	selection, s-possible nucleotide insertion)
14870	45238	A	14957	434	1790	GNFFRKTISACIISVLSYLVSCLR
						GASPTEASMDFCLKTSKREAAR
		ļ				ASRKICFVCKKKGAAINCQKDQ
						CLRNFHLPCGQERGLPFTIFWR
						VQIIL*QTSP/EHRTSNMGHVGE
						ESCILCCEDLSPTECLRTFQSPC
]	CSQAIYHRKCIQ/TNMPTHQSK
		ŀ				HFFKCPQCNNRKEFPQEM/REN
						GNSYSRQRCCLGTRARGFLRLI
		ŀ				SALSAL*CPHLVCMNKAETALR
						MKGGGASFCVLHADPTEPTGT
		1				APLLDLTVRNGSVRSVHL/PAA
						TDYIPENSGDIPCCSSTFHPEEHF
						CRDNTLEENPGLSWTDWPEPSL
						LEKPESSRGRRS/VLLEVQGCQ
		İ				NH*QLQKIQVTPSE*LLVPHNR
						V*TCAPPSGLGREHSGTVRQGS
		ļ				RASSETMS/LSKEKSQWSMRRE
		1				QSRCQQGNAFMATRVPLLSPPL
		ļ				LPPRILPP*SCFHMPLLTSPMFV
						VMQIKVFSPKKKKK
14871	45239	Α	14958	146	484	
14872	45240	Α	14959	228	353	
14873	45241	Α	14960	214	722	TARELPGRVLKEMYIQQK*EGA
						ALGRPTRPGRQGQRVRRAECR
		<u> </u>				RLFRGVLRPCNRC
14874	45242	В	14961	1	951	
14875	45243	Α	14962	241	517	QSCGQRLPTVLR/PLRTPWILPM
•						YPQPLPAGGPTPCLRSRPYINITI
				ļ		LKGDKGDPGPMGLPGYMGRE
						GPQGEPGP*GSKGDKGEMGRP
						RRPVP
14876	45244	Α	14963	748	1055	DRKQTSPGDAYHTIPGSLSSSSD
						HVLSEPD*GEGSHL**SVQHDD
						VPPTA**RLQY\RAVVLFPTATL
						PAILIM*GMRCPLVPRNCSRTW
						TAQIRAYIEVQQT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14877	45245	Α	14964	1	832	MPHVCAQRRLPVAAVLPPDGA
						QNCILLAGLSAGTFCHHGGVHI
						LLSNPVATSLLVAIDRVGQIPIPP
		ŀ				PPPPALPYDCCCPPPSRHTEDRE
						HVPIPLHWKLMVNRWFLPVPT
			:	:		EWIATLSNSSRSSRTCSMQGLSF
			;			LDSQVHLKAMIEDLRGNVSKES
						SPCLADNADVGQFPMEGGQKD
						PKQRIIPFCQGVTNLSGPHGPGA
						VCGGSHYQKQESSEISLSVGQV
						WTSSRRMSQVGGSSHCRGARL
						GPCNVPRRPDGVQDEFSLQPEE
						DAG*QTSVDPMVLEQYVVVA/
						NYQKQESSEISLSVGQ/VVDIIE
			:			KNESGWWFV/STAEEQGWVPA
						TCLEGQMGCRMSFLCSLKKML
						DWIQRRVSL
14878	45246	Α	14965	1	346	
14879	45247	Α	14966	157	413	ALVCS/SSLAIREMQIKTTMRYH
			į			LTPVRMAIIKKSGNNRCWRGC
						GEIGTLLHCWLDCKLVQPLWK
		<u> </u>				SVW*FLRNLELEIPFDPAIPLL
14880	45248	Α	14967	1930	5781	RAKSPANIIMTGSNSHITILTLN
						VNGLNSPIKRHRLASWIKSQDP
						SVCCIQETHLMCRDTHRLKIKG
						WRKIYQANGKQKKAGVAILVS
						DKTDFKPTKIKRDKEGHYMMV
						KGSIQQEELTILNMYAPNTGAP
				:		RFIKQVLSDLQRDLDSHTLIMG
						DFNTPLSTLDRSTRQKVNKDTQ
						ELNSALHQADLIDIYRTLHPKST
						EYTFFSAPHHSYSKIDHILGSEA
						LLSKCKRTEIITNYLSDHSAIKL
						ELRIKNLTQSR
14881	45249	A	14968	1	2235	
14882	45250	Α	14969	1	2148	
14883	45251	Α	14970	1	1821	
14884	45252	Α	14971	1	1248	
14885	45253	В	14972	4	2121	
14886	45254	Α	14973	1	1137	

NO:			D 2 Q	Nucleotide	Nucleotide location of fast	Amino acid sequence (X=Unknown,
	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14887	45255	Α	14974	1	1240	MNIDAKILNKILANRIQQHIKKL
						IHHDQVGFIPGMQGWFNICKSI
						NVIQHINRAKDKNHMIISIDAEK
						AFDKIQQPFMLKTLNKLGIDGM
						YFKIIRAIYDKPTANIILNGQKLE
						AFPLKTGTRQGCPLSPLLFSIVL
						EVLARAIRQEKEIKGIQLGKKE
						VKLSLFADDMIVYLENPIVSAQ
						NLLKLISNFSKVSGYKINVQKS
						QAFLYTNNRQTESQIMSELPFTI
						ASKRLKYLGIQLTRDVKDLFKE
						NYKPLLKEIKEDTNKWKNIPCS
						WVGRINIMKMAILPKVIYRFNA
						IPIKLPMPFFTELEKTTLKFIWNE
						KRARIAKSIPSQKNKAGGITLPD
						FKLYYKATVTKTAWYWYQNR
						DIDQWNRTEPSEITPHIYNYLIF
			-			DKPEKNKQIQKAFILH\KEPWV
						MWNMGTNVR
	45256	Α	14975	1	2241	
14889	45257	Α	14976	1	2802	MYGTGYRDVAGKWAIDPDSK
						REFLDVTQEGIQGCDFSGTCRQ
						TLSILTQPLRQWGLEGIKKPNS
						WIISEESVSNGGPPLLIPRQTAS
						GVDLQQTPTDLQLRVLTVRRK
						TNKQKGIASTSTKRTSTPKPHL
						YVTIIKDQSYIKPQRWGKNIAE
						KLKILKIRVALSLQRNAAPHQQ
						WNKAGRRMSLMSSQKKASEVI
						ESQMNEIKGEEKFREKRVKRNE
						QSLQEIWDYVKRPDLRLIGVPD
14890	45258	A	14977	1	2739	SDGENGTRLENTLQDI
	45259	A	14977	1	2478	
	45260	A	14979	590	727	
	45261	A	14980	1	3255	
	45262	A	14981	1	3810	
	45263	В	14982	1	3127	
		A	14983	1	3325	
		В	14984	1	1734	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14898	45266	Α	14985	11	2067	MELKTKARELREECRSLRSRCD
		ŀ				QLEERVSAMEDEMNEMKREG
						KFREKRIKRNEQSLQEIWDYVK
						RPNLHLIGVPETDGENGTKLEN
				1		TLQDIIQENFPNLARQANVQIQE
;						IQRTPQRYSSRRATPRHIIVRFT
						KVEMKEKMLRAAREKEIQTTIR
						EYYKHLHANKLENLEEMDKFL
			1			DTYTLPRLNQEKVESLNRPITG
		ļ				AEIVAIIN/SLPTKKSPGPDGFTA
		ŀ				EFYQRG\ILPNSFYEASIILIPKLG
						RDTAKKENFRPISLMNIDAKILN
						KILAKRIQQHIKKLIHHDQVGFI
						PGMQGWFNIRKSINVIQHVNRT
						KDKNHMIISIDEEKAFDKIQQPF
						MLKTLNKLGIDGMYFIIIRAIYD
				ŀ		KPTAHIILNGKKLEAFPLKTGM
						RQGCPLSPLLFNIVFKVLARAIR
						QEKEIKGIQLGKEEFKLSLFADD
			1	ļ		MIVYLENPIVSAQNLLKLISNFS
						KVSGYKINVQKSQTFLYTNNR
						QTESQIMSELPFTIASKRIKYLGI
						QLTRDVKDLFKENYKPLLKEIK
						EDTNKWKNIPCSWVGRINIVK
		ŀ				MAILPKVIYRFNAIPIKVPMTFF
		ŀ				TELEKTTLKFIWNQKRACIAKSI
						LSQKNKAGGITLPDFKLYYKAT
					:	VTKTAWYWYQNRDIDQWNRT
						EPSEITPHIYNYLIFDKPEKNKQ
						WGKDSLFNKWCWENWLAICR
						KLKLDPFLTPYTKINSRWIKD
14899	45267	Α_	14986	1	1293	
14900	45268	A	14987	1	3229	
14901	45269	Α	14988	1	3352	
14902	45270	Α	14989	1	1419	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14903	45271	A	14990	1	1716	MKAEIKMFFETNENKDTTYQN
!						LWDTFKAVCRGKFIALNAHNR
						KQERSKIDTLTSQLKELEKQEQ
						TYSKASRRQEITKIRAELKEIET
	}				İ	QKTLQKINESRSWFFERINKIDR
						PLARLIKKKRQKNQMDTIKND
						KGDITTDPTEIQTTVREYYKHL
						YTNKLENLEEMDKFLDTYTLPR
						LNQEEVESLNRPITGAEIVAIINS
						LPTKKSPGPDGFTAEFYQRYKE
						ELPGRDTTKEENFRPISLMNIDA
						KILNKILANQIQQHIKKIIHHDQ
						VGFIPGMQGWFNIRKSINVIQHI
	i					NRAKDKNHMIMSIDAEKAFDKI
						QQPFMLKTLNKLVLEVLARAIR
}						REKEIKGIQLGKEEVKLSLFAD
				- 		HMIVYLENPIVSAQNLLKLISNF
						SKVSGYKINVQKSQAFLYTNNR
	1					QTESQIMSELPFTIASKRIKYLGI
						QLTRDVKDLFRENYKPLLKEIK
						EDTNKWKNIPCSWVGRINIVKV
						AILPKVIYSFNAIPIKIPMPFFTEL
						EKTTLKFIWNQKRARIAKSILSQ
						KNKAGAITLPDFKLYYKPTVTK
						TAWYSYQNRD\IDQWNRTEPSE
						ITPHIYNYLIF
14904	45272	Α	14991	1	702	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14905	45273	A	14992	473	2385	HSFLSFIVLMYLTMDLTRSQKH GWASSPSHGTSPWPVRLRTSLA KPCKWQVSLCWWLSASPGSSF SLVRSSLLIIVPHAESVSRRPSLL SRVGLFPARY*HRVWQRSPSPQ TRDHLIKRNPSQDV*RRQQPRG TRVSNKCSLHTSALGIGRQPSC DATL*GQQGLP/VK*KRQQPAP/ MSQQQTGAALEVLARAIRQEK EIKGIQLGKEEVKLSLFADDMI VYLENPTVSAQNLLKLISNFSK VSGYKINVQKSQAFLYTNNRQ TESQIMSELPFTIASKRITYLGIQ LTRDAKDLFKENYKPLLKEIKE DTNKWKNIPCSWVGRINIVKM AILPKEPGWARSLVFGARGLQG QGRCGQSRGGGGSCLGGGGST ENWQHREPSARKTGSTENRRH REPAAPRTGSTENQHREPAAPR TGTENRHREPSARKTGSTENRR HREPAAPRTGSTENQHREPAAP RTGTENRHQEPAAPRTGGTENR QHREPAAPRTGSTENRQHREPAP RTGSTENRQHREPAPRTGSTEN RQHRESAAPRTGSTENRHQEPA APRTGTENRHREPSAPRTVGTE NRRHRELVAPRTGGTENRPREP PAGTRPRHTPMRGGSEGALFH

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	i	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
14906	45274	Α	14993	687	2274	LRDCKRINQHPVKTDQSAFCK
						M/API/MQDVVLEVLARAIRQEK
						EIKGIQLGKEEVKLSLFADDMI
					:	VYLENPIVSAQNLLKLISNFSKV
•						SGYKINAQKSQAFLYTNNRQTE
						SQIMSELPFTIASKRIKYLGIQLT
						RDVKDLFKENYKPLLKEITEDT
						NKWKNIPCSWVGRINIVKMAIL
						PKVIYRFNAIPIKLPMTFFTELE
						KTTLKFIWNQKRARMAKSILSQ
						KNKAGGITLPDFKLYYKATVT
1						KTAWYWYQNRDVDQWNRTEP
						SEITPHIYNYLIFDKPDKNKQW
						GKDSLFNKWGWENWLAICRKL
						KLDPFLTPYTRINSRWIKDLNV
						RPKTIKTLEENLDITIQDIGMGK
				•		DFMSKTPKAMATKAKIDKWDL
						IKLKSFCIAKETTIRVNRQPTKW
		}				EKIFATYSPDKGLISRIYNELKQI
		Ì				YKKKTNNPIKKWAKDMNRHFS
		ļ				KEDIYAAKKHMKRCSSSLAIRE
İ					į	IQIKTTMRYHLTPVRMAIIKKSG
						NN/R/CF*WQKPGPSG*ALRSWK
					ļ	EDIAGTSYVGMNQCQMRKNTK
						TLI
14907	45275	Α	14994	1	3189	
14908	45276	Α	14995	1	3351	MGKKQNRKTGNSKMQSASPPP
						KERSSSPATDQSWMENDFDEL
İ		İ				REEGFRRSNYSELREDIQTKGK
						EVENFEKNLEECITRITNTEKCL
						KELMELKTKARELREECRSLRS
						RCDQLEERVSAMEDEMNEMK
		İ				QEGKIKRDKEGHYIMVKGSIQQ
				•		EELTILNIYAPNTGAPRFIKQVL
		ľ				SDLQRDLDSHTLIMGDFNTSLL
						TLDRSMRQKVNKDTQELNSSL
			<u> </u>			HQADLIDIYRTLHPKSIEYTFFS
						APHHTYSKIDHIVGS
14909	45277	Α	14996	1	3192	
14910	45278	Α	14997	1	2742	
14911	45279	В	14998	1	2313	
14912	45280	Α	14999	1	1542	
14913	45281	Α	15000	1	3300	
14914	45282	Α	15001	1	1824	
14915	45283	A	15002		2982	

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14916	45284	Α	15003	1	2375	MSGFLISPKHKHGTNSGIVSWA
						DVQSQLRHIEAAFDQEAAAVL
						MARLGVFRAESEEGPDVLRWL
						DRQLIRLHSLLTTLDIVPAGKGE
						VVKGPMSISQQAKGCQKFGQY
		ŀ				NKEDPTSFRLSDSFSLYPQFMF
						HLRRSPFLQVFNNSPDESSYYR
						HHFARQDLTQSLIMIQPILYSYS
						FHGPPEPVLLDSSSILADRILLM
						DTFFQIVIYLGETIAQWRKAGY
						QDMPEYENFKHLLQAPLDDAQ
		ļ				EILQARFPMPRYINTEHGGSQK
				i		KREKNQIDAIKNDKGDITTEPTE
1						IQTTIREYYKHLYANKLESLEE
1						MNKFLDTYTLPRLNQEEVESLN
						RQITGSEIEAIINSLPTKKSPGPD
<u> </u>						RFTAKFYQRTNDQKHMIISIDA
					ŀ	EKAFDKIQQPFMLKTLNKLGID
						GTYFKIIRAIYDKPTANIILNGQ
						KLEAFPLKTGTRQGCPLSLLLF
						NIMLEVLARAIRQEKEIKGIQFG
	1					KEEVKLSLFADDMIVYLENPIV
						LAQNLLKLISNFSKVSGYKINV
						QKSQAFLYTSNRQTESQIMSEL
						PFTIASKRIKYLGIQLTRDVKDL
		İ				FKENYKPLLNEIKEDTNKWKNI
						PCSWVGRINIVKMAILPKVIYRF
						NAIPIKLSMTFFTELEKTTLEFI
		1			1	WNQKRARITKSILSQKNKAEGI
						TLPDFKLQYKATVTKTAWYSY
						QNRDIDQWNRTEPSEIVPCIYN
						YLIFDKPDKNKKWGKDSLFNK
14917	45285	Α	15004	1	1428	
14918	45286	Α	15005	2	1975	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14919	45287	A	15006	1	3891	MKPEIKMFFETNENKDTTYONL
						WDTFKAVCRGKFIALNAHKRK
						QERSKINTLTSQFKELEKQEQT
1						NSKASRRQEITKIRAELKEIETQ
						KPLQKINESRSWFFEKINKIDRP
						LARLIKKKREKNQIDAIKNDKG
						DITTDPTEIQTTIREYYKHLYAN
						KLENLEEMDKFLDMYTLPRLN
						QEEVESLNRPITGSEIEAIINSLP
						TKKSPGPDGFTAEFYQRYKEEL
ŀ						VSFLLKLFQSIKKEGILLNLFYE
						DSIILKPKPGRDTTKKENFRPISL
						MNIDAKILIKILANRIQQHIKKLI
						HHDQVGFIPGMQGWFNIRKSIN
İ						VIQHINRTKDKNHMIISIDAEKA
						FDKIQQPFMLKTLNKLGIHGTY
						LKIIRAIYDKPTANIILNGQKLK
						AFPLKTGTRQGCPLSPLVFNIVL
						EVLAWAIRQEKEIKGIQLGKEE
						VKLSLFADDMTVYLENPIISAQ
						NLLKLISNFSKVSGYKINVQKS
			1			QAFLYTNNGQTESLIMSELPFTI
						ASKRIKYLGIQLTRDVKDLFKE
1						NYKPLLNEIKED\TNKWKNIPCS
						WVGRINIVKMAILPKVIYRFNAI
ŀ						PIKLPMTFFTDLEKTTLNFIWNQ
						KRARITKSILSQKNKAGGITLPD
						FKLYYTATVTKTAWYWYQNS
]	MVLVPKQRYRSMEQNRALRN
						NAACLQLSDL
14920	45288	Α	15007	1	4215	MHSCLPDATGEGELYNSLGSLT
14920	43200	A	13007	1	4213	WSFLSMIGVLLWVDFSGDSIDL
						CSPLWNRTNLEALQKKLEELEL
	j					DEQQRKRLEAFLTQKQKVGEL
				ŀ		KDDDFEKISELGAGNGGVVFK
						VSHKPSGLVMARKLIHLEIKPAI
						RNQIIRELQVLHECNSPYIVGFY
						GAFYSDGEISICMEHMVLLQER
						PCPGFLWMENSDYGLLLPLDLL
	:		1			LCLSFSFSKVLSDLVSDILIHGM
						KARNRSIHGDVVVVELLPKNE
1.10=	1.5000	ļ	1.5000		2000	WKGRTVALCENDC
14921	45289	Α	15008	1	2823	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14922	45290	A	15009	1	3253	 MGDFNTPLSTLDRSTRQKVNK
17722	43270	'`	1.500)	1	5233	DTQELNSALHQGDLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKWKRTEIITNYLSDH
			}			SAIKLELRIKNLTQSRSTTWKLN
ļ			ļ			NLLINDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFIALNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQKTLQKINESR
						SWFFERINKIDRPLARLIKKKRE
					:	KNQIDTIKNDK
14923	45291	A	15010	1	2816	MEDEMNEMKREGKFREKRIKR
11,723	13271	`	1.5010	1		NEQSLQEIWDYVKRPNLHLIGV
į		}				PESDVENGTKLENTLQDIIQENF
		ł				PNLARQANVQIQEIQRTPQRYS
						SRRATPRHIIVRFTKVEMKEKM
						LRAAREKGHCNSGKSHRSLWE
						QNKGIFTTCICDDAEDLQSPSTA
						GPWFGVLYTKGGPYPQEDIFSE
		1				EVHTGPKLRKKIQEYQLTSKWS
						KSDVQVSVERRMAGGNPNQCH
						IGEVLLDGFTAEFYRRYKEELV
ļ						PFLLKLFQSIEKEG
14924	45292	A	15011	1	3347	MGDFNIPLSTLDRSTRQKVNKD
						TQELNSALHQADLIDSYRTLHP
		•				KSTEYTFFSAPHHTYSKIDHILG
				1		SKALLSKCKRTEIITNYLSDHSA
						IKLELRIKNLTQNCSTTWKLNN
						LLLNDYWVHNEMKAEIKMFFE
						TNENKDTTYHNLWDAFKAVCR
						GKFIALNAHKRKQERSKIDTLT
						SQLKELEKQEQTHSKASRRQEI
		ŀ				TKIRAELKEIETQKTLQKINESR
						SWVFERINKIDRPLARLTKKKS
		<u> </u>				EKNQIDAIKNDK
14925	45293	Α	15012	1	3855	MGKKQNRKTGNSKTQSASPPP
						KERSSSPATEQSWMENDFDELR
						EEGFRRSNYSELWEDIQTKGKE
						VENFEKNLEECITRITNTEKCLK
		1				ELMELKTKARELHEECRSLRSR
		1				CDQLEERVSAMEDEMNEMKRE
						GKFREKRIKRNEQSLQEIWDYV
						KRPNLRLIGVPESDVENGTKLE
						NTLQDIIQENFPNLARQANVQI
						QEIQRMPQRYSSRRATPRHIIVR
						FTKVEMKEKMLRAARQKAPH
14006	45204	<u> </u>	15012	1	2550	HTYSKIDHIVGSKAL
14926	45294	A	15013	1	2559	
14927	45295	A	15014	1	2956	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14928	45296	A	15015		1853	MGRNQSRKAENSKNQSASSPP KDCSSSPAMEQSWTENDFDELT EVGFRRSVITNFSELKEHVETNL KEAKHLEKRLDKWLTRIKSVE KILNDLMELKTMAQELHDTCR SFNSQFNQVEERVSVIEDQINE MNIDGTYLKIIRAIYDK\PPRNII LNQQKLEAFPLKTGTRQGCPLS PLLFNIVLEVLARAIRQEKEIKGI QLGKEEVKLSLFADDMIVYLEN PIVSAQNLLKLISNFSKVSGYKI NVQKSQAFLYTNNRQTESQIMS ELPFTIASKRIKYLGIQLTRDVK DLFKENYKPLLKEIKEDTNKW KNIPCSWVGRINIVKMAILPKVI YRFNAIPIKLPMTFFTELEKTTL KFIWNQKRARIAKSILSQKNKS GGITLPDFKLYYKATVTKTA/W L/YHTPLSCLPP*WNRTEPSEIIP HIMAVAQIQQVKLLLLQGQHA SSRLQPTSSTKCGGWTASSASP RSMGLSSWKGNLASAIHYFPCK PLTLPSRISTSSSSWGCRQAHAV LEVLRWPPGLGWPGWHHLTLL HVPPVRITKSNGPRACTRASAS
						HARHHQYNQEPSLSPLHSPLPP PEQVLVSTAERTEDRSYHRTLC
						RHSPYFKNEAADLRGEGYSS
14929	45297	В	15016	1	3276	
14930	45298	Α	15017	1	2274	
14931	45299	В	15018	440	3383	
14932	45300	Α	15019	1	2646	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		1
	sequence	:	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14933	45301	A	15020	1	2229	 MGKKQNRKTGNSKTQSASPPP
						KERSSSPATEQSWMENDFDELR
						EEGFRRSNYSELREDIQTKGKE
		ľ				VQNFEKNLEECITRITNTEKCLK
			i			ELMELKTKARELREECRSLRSR
						CDQLEERVSAMEDEMNEMKRE
		ŀ				GKFREKRIKRNEQSLQEIWDYV
						KRPNLRLIGVPESDVENGTKLE
			i .			NTLQDIIQENFPNLARQANVQI
						QEIQRTPQRYSSRRATPRHIIVR
			;			FTKVEMKEKMLRAAREKEIQT
						TIREYYKHLYANKLENLEEMD
						TFLDTYTLPRLNQEEVESLNRPI
		İ				TGSEIVAHNSLPTKKSPGPDGFT
						AEFYQRYKEELVPFLLKLFQSIE
						KEGILPNSFYEASIILIPKPGRDT
						TKKENFRPISLMNIDAKILNKIL
						ANRIQQHIKKLIHHDQVGFIPG
						MQGWFNIRKSINVIQHINRAKD
						KNHMIISIDAEKAFDKIQQPFML
			•			KTLNKLGIDGTYFKIIRAIYDKP
						TANIILNGQKLEAFPLKTGTRQ
		l				GCPLSPLLFNIVLEVLARAIRQE
						KEIKGIQLGKEEVKLSLFADDM
			1			IVYLENPIVSAQNLLKLISNFSK
						VSGYKINVQKSQAFLYTNNRQ
						TESQIMGELPFTIASKRIKYLGIQ
						LIRDVKDLFKENYKPLLKEIKE
						DTNKWKNIPCSWVGRINIVKM
]	AILPK\IGKTTLKFIWNKKSRIAK
						SILSQKN/KAGGITLLTQLYYKA
						TVTKTAWYWYQNRDIDQWNR
14934	45302	Α	15021	1	960	
14935	45303	A	15022	1	2928	MEDEMNEMKQEEKFREKRIKR
						NEQTLQEIWDYVKRPNLHLIGV
		ŀ				PESDGENGTKLENTLQDIIQENF
						PNLARQANVQIQEIQRMPQRYS
						SRRATPRHIIVRFTKVEMKEKM
						LRAAREKDRSMRQKVNKDTQE
						LNSALHQADLIDIYRTLHPKSTE
				i		YTFFSAPHHTYTKIDHILGSKAL
						LRKCKRTEIITNYLSDHSAIKLE
						LRIKNLTQNRSTTWKLNNLLLN
						DYWVHNKMKAEIKMFFETNEN
						KDTTYQNLWDAF
14936	45304	A	15023	1	1782	TO THE VIEW DIE
14937	45305	A	15023	1	4881	
14937	45306	B B	15025	1	5962	
14730	1-3300	ا	13023		3702	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14939	45307	Α	15026	1	2053	MKAEIKMFFETNKNKDTTYQN
14757	45507	A	13020		2033	LWDAFKAVCRGKFIALNAHKR
						KQERSKIDTLTSQLKELEKQEQ
						THSKASRRQEITKIRAELKEIET
						QKTLQKINESRSWFFERINKIDR
						1
						PLARLLKKKREKNQIDAIKNDK
						GDITTDPTEIQTTIREYYKHLYA
						NKLENLEEMDKFLDTYTLPRLN
						QEEVESLNRPITGAEIVAIINSLP
						TKK\SPGPDGFTAELYQRYKEG
						AVLEVLARAIRQEKEIKGIQLG
						KEEVKLSLFAGDMIVYLENPIV
			,			SAQNLLKLISNFSKVSGYKINV
						QKSQAFLYTNNRQTESQIMSEL
1						PFTIASKRIKYLGIQLTRDVKDL
Ì						FKENYKPLLNEIKEDTNKWKNI
1						PCSWVGRINIVKMAILPKVIYGF
						NAIPIKLPMTFFTELEK\TT*KFI
ļ						WNQKRARIAK*ILSQKNKAGGI
				:	:	TLPDFKLYYKATVTKTAWYW
						YQNRDIDQWSRTEPSEIMPHIY
						NYLIFDKPDKNKKWGKDSLFN
						KWCRENWLAICRKLKLDPFLTP
						YTKINSRWIKDLNVRPKTITTLE
İ						EKLSNTIQDIGMGKDSMSKTPK
						AMATNVKIQKWDLIKLKSFCT
				ļ		AKETTIRVNRQPTTWEKIFATY
						SSDKGLISRIYNELKQIYKKKTN
l						NPIKKWVKDMNRHFSKEDIYA
						AKKHMKKCSPSLAIREMQIKTT
-						MRYHLTPVRMAIIKKSGNNRC
14940	45308	Α	15027	3	3229	
14941	45309	A	15028	1	3907	MGDFNTPLSTLDRATRQKVNK
			12020	1		DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIL
		ľ				GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQNRSITWKLSN
						LLLNDYWVHNEMKADIKMFFE
						TNENKDTTYQNLWDTFKAVCR
					1	GKFLALNAHKKKQERSKIDTLT
				ŀ		SQLKELEKQEQTHSKASRRQEI
						TKIGAELKEIETQKTLQKINESR
	1					SWFFERINKIDKPLARLIKKKRE
14042	45210	_	15020	1	1005	KNQIDAIKNDK
14942	45310	A	15029	287	4005	
14943	45311	A	15030		2708	
14944	45312	A	15031	1	4602	
14945	45313	Α	15032	1	4806	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14946	145214	1	15022	965	5928	 TWKGTTSTSRCKIMPKYRSTRQ
14940	45314	Α	15033	903	3928	1
•						KVNKDTQELNSALHQADLIDIY
						RTLHTKSTEY/TFFSAPHHTYSK
		1				IDHIVGSKALLSKCKRTEIITNY
		1				LSDHSAIKLELRIKNLNQSRSTT
						WKLNNLLLNDYWVHNEMKAE
						IKMFFETNENKDTTYQNLWDA
						FKAVCRGKFIALNAHKRKQERS
						KIDTLTSQLKELEKQEQTHSKA
		1				SRRQEITKIRAELKEIETQKTLQ
						KINESRSWFFERINKIDRPLARLI
		<u> </u>				KKKREENQID
14947	45315	Α	15034	383	8708	
14948	45316	Α	15035	2	3257	WRKIYQANGKQKKAGVAILV/S
		1				DKTDFKPTKIKRDKEGHYIMVK
						GSIQQEELTILNIYAPNTGAPRFI
İ						KQVLSDLQRDLDSHTLIMGDFN
						TPLSTLDRLMRQKVNKDTQEL
		ŀ				NSALHQVDLIDIYRTLHPKSTE
		ł				YTFFSAPHHTYSKIDHILGSKAL
						LSKCKRTEIITNYLSDHSAIKLE
						LRIKNLTQSRSTTWKLNNLLLN
		1				DYWVHNEMKAEIKMFFETNEN
						KDTTYQNLWDAFKAVCRGKFI
						ALNAYKRKQERS
14949	45317	Α	15036	1	3508	MELKTKARELREECRSLRSPCN
						QLEERVSAMEDEMNEMKREG
ŀ						KFRDKRIKRNEQSLQEIWDFVK
ŀ						RPNLRLIGVPESDGENGTKLEN
İ						TLQDIIQENFPNLARQANIQIQEI
ľ		ŀ				ORTPORYSSRRATPRHIIVRFTK
ľ						VEMKEKMLRAAREKAPHHTYS
		ł				KIDHILGSKVLLSKCKRTEIITN
						YLSDHSAIKLELRIKNLTQNHST
		1				TWKLNSLLLNDYWVHNEMKA
						EIKMFFETNENKDTTYQNLWD
						TFKAVCRGKFIALN
14950	45318	В	15037	1	4760	
14951	45319	A	15038	1	3110	
14952	45320	Α	15039	1	5546	MELKTKARELREECRSLRSRRN
			[QLEERVSAMEDEMNEMKREG
						KFREKRIKRNEQSLQEIWDYVK
						RPNLRLIGVPESDAENGTKLEN
						TLQDIIQEDFPNLARQANVQIQE
						IQRTPQRYSSRRATPRHIIVRFT
			Ì			KVEMKQKMLRAAREKDFKPT
						KIKRDKEGHYIMVKGSIQQEEL
						TILNIYAPNTGAPRFIKQVLSDL
						QRDLDSHTLIMGDFNTPLSTLD
	1			•		RSTRQKVNKDTQELNSALHQA
						DLIDIYRTLHPKSTE
L	<u> </u>		L	L	L	DEIDITKIERTKSIE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14953	45321	В	15040	1	4534	
14954	45322	В	15041	1	1826	
14955	45323	В	15042	1	1626	
14956	45324	A	15043	248	1212	NSKHYRSGQKKKLPTLPRAAN SRWRSLKI/GFTVSFDETLKPFV RDVSGSRLKDLPKPNKSDDETR ANDAVNRYKLLKKDARTIAAQ QVARLESAMCLRRWSLENFQ LFLVEHPLVRHLTRRLIWGVYS AENQLLACFRVAEDNSSSTADD DLFTLPEGDISIGTPHVLEISPTD AAALLFADYELLPPFRQLDRNS YALTEAERNASELTRWAGRKC PSGRVMGLANKGWIKGEPQDG GWIGWMIKPLGRWSLIMEIDEG FAVGMSPAELSAEQLLSKLWL WEGKAERYGWGSNSTQEAQFS VIDAITASELINDIEALFE
14057	45225	_	15044	1	1788	VIDATTASELINDIEALFE
14957	45325	A	15044 15045	1	3048	
14958	45326	A	15046	1	3546	
14959 14960	45327 45328	A A	15047	969	1205	SPVRCCISAVHRTYHRRTSALL
14900	43326	A	13047	909	1203	RLAELHRLLEKCASRQWMYEN I*SPFRQIA*CLPYSDHKNQRLH GLDSIFLARLLW
14961	45329	A	15048	877	1248	WPLTVPYCGVGYAS*A*VADN HRAVGYLSRWAGRKCPSGRV MG/LANKGWIKGEPQDGGWI/G WMIKPLGRWSLIMEI/DEGFAV GMSPAELSAEQ/LLSKLWLWEG KAERYG/WGSNSTQEAQFSVID
14962	45330	A	15049	854	1313	DHGVASAATCVYTNEGHRQSY VRVCEYASRTRVSHHCSV*RVL RCEAIRIVSPDRSC*RDPEFSYL\ ALPENYNRLFLPNSTNQTNNRI KTLNSIAIGKLLAAGGVYNGNI EGFRDTAEKLDGRTIDGYDQIL NEKTAVIAKATA\SILLTKRS
14963	45331	A	15050	1010	1396	KPFGKLTYKLFEGHAWLVKAE KVRVGYRILGRLTYPFDIRFPAL PFGLVL*KGAHFTVPRIWAGHA C*LMPMALPMFRWKVTGLLFIP ICLVKPSFLTLTIIIAIRRISTSTN CRKTPKQPSRWCKPR

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14964	45332	A	15051	361	837	DGQQLIALHRLALRELQQAVH AGLPQQAKILFDGGSEIGKIPSEI YFCQLVKLILCPVLFPCWRRVA ILWRRRDVLFFGSFSFSALVSPH LCG/SYLPFVFDVGDRL*LSVG WS\KEIWPCK*GFGHLMSVLNT PMKSLVEGFKHFPLAGFGYEYL SFFCR
14965	45333	Α	15052	1563	1652	
14966	45334	A	15053	2756	2995	AMRSAHRCTVFSALLNCWQITP HLTPSVMICGQLLTLVNRC*PSS TIFSIIPLSKQVARMFRSAMSPL NRARCWKVPCN
14967	45335	Α	15054	762	2184	WRAVPSEPISFILKADTEKYVY L*NSSSAQYSGSCTRAYYLDTQ HPTPRLCLIF\GGKDSGLMLHLT AELARQMGKKICVLFIDWEAQ FSCTINYVQSLRELYTDVIEEFY WDALPLTTQNSLSQYQPEWQC WEPDVEWVRQPPQDAITDPDFF CFYQPGMTFEQFVREFAEWFSQ KRPAAMMIGIRADESYNRFVAI ASLNKQRFADDKPWTTAAPGG HSWYIYPIYDWKVADIWTWYA NHQSLCNPLYNLMYQAGVPLR HMRICEPFGPEQRQGLWLYHVI EPDRWAAIGSPADREEDAEEYL EAIMEARVTVAGMGLVMEVQ DYFDGEADRLAKAWLAEYTPQ IKSLKDERKEAYRQIVEMSTEP QDVDLVRPANKFEMTRVREGE KEADLPVWKHHLLCDESGNYP ALLNHWETKVFEIETKREGFAF WYRNPQYTGQSSLGIAYVEAE QYKIVRPDFLFFAEQDGKMVV
14968	45336	A	15055	2069	2586	PWLENLPVGICWSAVKKMGGT RSLVSCL/AVTLVSPQLMNAYL LGQQLPEVWDFGMFSIAKVGY QAQVIPALYLVVVPVCSLILAV FLAHALIGPFGRMIGDGVAFAV RHLMTGSFAPIGAALFGFLYAP LVITGVHQTTLAIDLQMIQSIGC TPVLPVNALSNIDLWHISILV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
14969	45337	Α	15056	356	1205	APASMTFEQFVREFA/VMVFTK
						RPAAMMIGIRADESYTVLSPSP
						V*INNVLPTITWTFPQTIYDFLW
	i					LIGEAIFF/YLPVGICWSAVKKM
						GGTRSLVSCL/AVTLVSPQL/MN
						AYLLGQQLPEVWDFG\VCSLIL/
						AVFLAHALIGPFGRMI/GDGVA
						FAVRHLMTGSFA/PIGAALFGFL
						YAPLVIT/GVHQTTLAIDLLMM
						QSMGGTPVWPLIALSNIAQGSA
						VIGIIISSRKHNEREIPGEVFPRT
	İ					GVGKGKSLG*PLH*LQPYRQHL
						GAGSYSQTLKKQIAESAVRLSIP
14970	45338	В	15057	1	435	
14971	45339	Α	15058	2	1785	WLLYRAHVKGE/STEALLPNM
	1					VATSLAKLPIPKLMRWGATYV
	ŀ					HFVRPVHTVTLLLGDKVIPATIL
	ŀ					GIQSDRVIRGHRFMGEPEFTIDN
	ł			1		ADQYPEILRDSGKVIADYEERK
	ļ					AKIKADAEEAARKIGGNADLSE
						TLLEEVASLVEWPVVL/TAKF/E
						EKF/LAISRTALVPYSADNMYQ
						LVNDVQSYPQFLPGCTGSRILES
						TPGQMTAAVDVSKAGISKTFTT
						RNQLTSNQSILMNLVDGPFKKL
						IGGWKFTPLSQEACRIEFHLDFE
	ŀ					FTNKLIELAFGRVFKELAANMV
						QAFTVRAKEVYSASMRYIYVIV
						YISYRPPNLLDSHIIGFSSITTHC
						AVTSHHYNSCDMTMTIHSYVT
			Ì			PYISVRVSLTQTYAPSPSRNLRI
						TDLTIYRAISSEKSIDDTRHIHTI
		į				ESRCRVCGAHDSRDAERLDRF
						AQTADEIRIADRGFSSRPECIRS
						LAFGEADYIVRVHWRGLRWLT
						AEGMRFDMMGFLRGLDCALIS
						KTRLLSENRRKGRVVQAETLE
						AAGHVLLLTSLPEDEYSAEQVA
						DCYRLRWQIELAFKRLKSLLHL
						DALRAKEPELAKVGIFANLLAA
						FLIDDIIQPSLDFPPRSAGSEKKN
14072	45240	 	15050		002	FLIDDIIQFSLDFFFKSAUSEKKN
14972	45340	A	15059]]	993	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	,	
14973	45341	Α	15060	1	5621	MSEKTFLVEIGTEELPPKALRSL
						AESFAANFTAELDNAGLAHGT
						VQWFAAPRRLALKVANLAEAQ
						PDREIEKRGPAIAQAFDAEGKPS
		l				KAAEGWARGCGITVDQAERLT
						TDKGEWLLYRAHVKG\EGTEA
						LLPNMVASSVAKLPIPKLMRW
						GASDVHFVRPVHTVTLLLGDK
						VIPATILGIQSDRVIRGHRFMGE
				ŀ		PEFTIDNADQYPEILRERGKVIA
						DYEERKAKIKADAEEAARKIGA
						KAWGTNVEFHVQNQP
14974	45342	Α	15061	82	732	
14975	45343	Α	15062	1	3849	MDLGANGWQTFRYVVLPNLSS
		ŀ				ALLAGGMLAFALSFDEIIVTTFT
						AEQVDAAVRAADAAFAEWGQ
						TTPKVRAECLLKLADVIEENGQ
						VFAELESSNNTAFARASSNGDL
						PTKADLQAQLDSLNKQKDLSA
	1					QDKLVQQDLTDTLATLDKIDRI
	1					KEETVQLRQKVAEAPEKMRQA
						TAALTALSDVDNDEETRKILST
						LSLRQLETRVAQALDDLQNAQ
		İ				NDLASYNSQLVSLQTQPERVQ
14076	45244	ļ.	15062	210	1186	NAMYNASQQLQQIRSRLDG
14976	45344	Α	15063	219	1180	PPKLGRKRGIFLPSHSNPNTNRI CPNCCMNGVSTANTTSKSSLVI
	Ì					CAWTKHALLNAILQPEQLWKS
	ŀ					GSRWRINLALPRGYRESLLQPW
						QAERLTTDKGEWLLY/RAHVK
						GESTEALLPNMV/ATSLAKLPIP
						KLMRWG/ASDVHFVRPVHTVT
						LL/LGDKVIPATILGIQSDRVIAA
						TGKAKIKADAEEAAR/KIGGNA
						DLSESLLEEV/ASLVEWPVVLN
						AKFDEK/FLAVPAEALVYTMKG
						DQ/KYFPVYANDGK/LLPNFI/F/
						VANIESKDP/Q/QIISGNEKSFVR
						LADAEFFF/NTT*KRLFLPHQKL
						PPSQILHQQTRNYASFCHRHRD
						NWTSA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
14977	45345	Α	15064	11	1496	SGGCAGLHGGLCSKEPFQASV
						QQGVAGHACAPAALQVPVCGT
						LSDCGSRAASLCENPSFISLYGR
						LVATSLAKLPIPKLMRWGASD
						VHFVRPVHTVTLL\LGDKVIPAT
						ILGIQSDRVIRGHRFMGEPEFTI
						DNADQYPEILRERGKVIADYEE
				-		RKAKIKADAEEAARKIGGNAD
				•		AKFEEKFLAVPAEALVYTMKG
						DQKYFPVYANDGKLLPNFIFVA
						NIESKDPQQI/DRIQALAGWIAE
						Q\IGADVNHATRAGLLSKCDLM
						TNMVFEFTDTQGVMGMHYAR
						HDGEAEDVAVALNEQYQPRFA
				1		GDDLPSNPVACALAIADKMDT
	1					LAGIFGIGQHPKGDKDPFALRR
1						AALGVLRIIVEKNLNLDLQTLT
						EEAVRLYGDKLTNANVVDDVI
1						DFMLGRFRAWYQDEGYTVDTI
-						QAVLARRPTRPADFDDWHALG
İ						RAGTGDVVVGVRGRLAIVPLT
						DGLDEAGHSTGVLQGEVPQRL
]		ERAATVPGVTLQPQALQARGL
14978	45346	A	15065	134	384	VTPTGWRKRRIRQWCSDAGCG
1770	13310	` `	15005			VNALSGLRFSHSLALFSAAMFI
1						LGVVSGITMSISP*ILSCVRSKNP
						LLVSGSGPLSCNSTIH
14979	45347	A	15066	11	723	TKSPKIARSRHIPLCHFIFERRPI
'''	1.55.7	`	1.000	<u> </u>	, 25	MTNSNRIKLTWISFLSYALTGA
						LVIVPGMVMENIADYFNLPVSS
				ĺ	•	MSNTFTFLNAGILISIFLNAWLM
						EIVPLKTQLRFGFLLMVLPVAG
	1					LMFSHSLALFSAAMFILGVVSGI
						TMSIGTFLVTQMYEGRQRGYR
	}	İ				LFLAASLFSMAG\MIFPMIAAFL
1		i				LAPSS*W*WVYACLGLAYVTIF
						IMTFGCEFPALGKHAPKSYAPV
						EKVKWGILGGASSFRAV
14980	45348	Α	15067	422	1045	LLEGKLTNRKQ*HQH*QKARPC
						KKPIQSAQN/WNREEYDELTEV
	ĺ					GRRWVIITNSFELKEHVLTQWK
						EAKNLDKRLQEGLTRITSLEKKI
						NDLMVLKNTARELHEAYTSINS
						RIDQAEERISEIEDQLNEIKLED
						KIRGKRIK/RQTNKQSL*KMWD
		1				YVKRPNLRLIGVPESDEENVTE
						LENTLQDIIQENFPNLARQANIQ
					;	IQEIQTLLR
14981	45349	A	15068	1	1131	
			L		1	l

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14982	45350	A	15069	383	756	LNPSNKLALINPAASPIIKGVDL TANQI/PCIADIPLYDADVQQE* CFPATVEALGEQIRQADGVVIV TPEYNYSVPGGLKNAIYWLSRL RQILVFLDAMVMNKPEFMGGV IQNKVDPQTGKVIDH
14983	45351	Α	15070	343	924	
14984	45352	A	15071	223	308	
14985	45353	A	15072	57	5546	NKMARKKFSGLEISLIVLFVIVT IIAIALIVVLATKTPAVDEISDST STPATTRVTTNPSDSGKCPNVL NDPVNVRINCIPEQFPTEGICAQ RGCCWRPWNDSLIPWCFFVDN HGYNVQDMTTTSIGVEAKLNRI PSPTLFGNDINSVLFTTQNQTPN RFRFKITDPNNRRYEVPHQYVK EFTGPTVSDTLYDVKVAQNPFS IQVIRKSNGKTLFDTSIGPLVYS DQYLQISARLPSDYIYGIGEQVH KRFRHDL
14986	45354	С	15073	1	1818	-
14987	45355	Α	15074	493	576	
14988	45356	Α	15075	381	498	
14989	45357	Α	15076	48	514	LLLFLLLQSYDWSESGKRQSAL HGRQKGLPKSPTAARGGSDPW PSCRHP*GPNRKRRKS/R*PMET APRRPSVSPALTPARPPGLTATP RLHPRPP/RSPNPSANLLRRPTP DT*AHATFGPGRSRHHFPGQLS TE/RRRSAEGGRSRRNEASRRE GGE
14990	45358	В	15077	1	1116	
14991	45359	С	15078	249	332	
14992	45360	A	15079	710	1326	RAAGNCFPEPGFVKQLSAAAA VVAPAAAATEGSIPDTGKSPSSI SHWPNPQKNVICKRRGRVEQE QAWEKGR*RVVFGL*KSMRCP VDLLPQSCLCQRFLHQ/GKWIL TKDYIIHSAKSGRWLDATTYE WGYKIEKDSRYSPQMQSAPKR WREELKRTGAPGAFHRWKVVL LVRTDKRSDSLIRFSDNFGDCDI RMEEKTFRTLMES

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
14993	45361	A	15080	602	1257	FGKRGRTITLCILAVGMLCCRK GHTKPRRRPGASPISTHLHSSPPI STHLHTSPPPHRREEAR\SGQ*R QSCGAAAAR*PWWWNCDGWG PKCQRFSKH*DIAQPDEVSWGQ *PWAVGALAGPHP\PQASEGPL GCTPASICGSTRKPSKPLSKKPP QAARTSSSRIVIFTPISTRPPTQG QPEKAKSATPTNDLEGPSAKVK FVNFLMDETFPEPSFFI
14994	45362	С	15081	140	175	
14995	45363	A	15082	2295	3089	NLLGYQEINPPLQIFWDKLLAV SERDSSVKEESGDVESDDTML* MLPCP*ETTEFSS*PFRISS*LSS* STAQI\PAETCFAGGWYSIFCPQ RIEQETMCLL*SWQSEVAVSCE SPWPTALHHSHPNSRAPPPCY/S KSTSSSLSSPLENSG*QNRPPSLL ITSVLPLRITCLP*VSFIPPIMQLG GSFLSNDRL*QSFILPKNVSLEF VSFPLI*GAVSKPAISFNTSVSPL
						LVSDIFFALSLLP*GLGTWLIPSS SSLLLVRGF
14996	45364	В	15083	143	1727	
14997	45365	В	15084	307	1017	
14998	45366	A	15085		959	MSRMGLLGFGERKELPNHPAV SEWPLTLVTGCSSGTVAEGEGN LTTSGLFSIVPLHPDGVAAGQR HSSLPRWGSQAEALLTSQTVDG QAEMLLTPQTMGGQEEMLLTS QTTGEDFITRSTESGTHSKSHKG DEGEPRWMLGWMLWVCDIAE PLVKAKLLDCKHTCCSVCLQQ TRTSQKDVRCPWCSGVTKLPPD FSVSQLLDDPEVLAVITIPHASE HMPVFIRLPGSGCY\SCPCPSPRS TRCCPEI*AATCCPGTGRSPSL\C DHPCWTAASARQGSPGGGGG AGQAGHGEKLHVVEGVHCHL GGLRLGLPPQHRAP
14999	45367	В	15086	112	672	
15000	45368	A	15087	2	530	SGRSAALLSARVNRSQSFAGVL GSHERGPRSFPVFSPPGPPRKPP ALSRVSRMFSVAHPAAKVPQPE RLDL\VYTALKRGPDGLLGAVH QQEQEKLQGQIRESKRNS/RLG EWREGFLYDLDKQVKSIERFLR RLEFHASKIDELYE\AYCVQ\RR LR\DGALHMG\RAYTTGSPGSR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15001	45369	A	15088	216	1369	EPVLRRRPRQPRAGAQGFPVFS
						PPGPPRKPPALSRVSRMFSVAH
						PAAKVPQPERLDLVYTALKRG
i						L/QAYLEVHQQEQ/DKLQGQIR
						ESKRNSRLVMAGGVLWTVV*E
						RGAVTSLSASAGVLLIPPKYLPL
						LPTPAGL\LYDLDKQVKSIERFL/
						RRLEFHASKVRVQHVCRWD\D
						ELYEAYCVQRRLRDGAYNMV
						RAYTTGSPGSREARDSLAEATR
						GHREYTE/DEGWGPMRQRHRS
						MCLLES/ELEAQLGEFHLRMKG
						CAVDINDLGTIKLSLEVTGAAS
		ŀ				SVNKASTVTKRSPPIARAHRTH
						PHFGNSFLSTRKPNTYTANHRR
]			RPIEQHPDMRTPPPPHRARTNT
			1			GRKDKVRAHEGEKATPTRTRG
			:			APHGKAQRLRPLYQYVTIATLS
15002	45370	Α	15089	1	940	MSYNSKNYVETITTMNSVKFG
15002	13370	` `	13003	 		ETTPRNRTHYLSTNEKKIFLSGV
						FSNAYPAGLSRGWDSPWGSTP
						WVRHFCYLSSTQLCDQNCSLLS
			<u> </u>			SLLPAGPSVCNRLLASAWQKSS
						SSASSEASETCQSVSECSSPTSD
						WSKVGSHEQPSGATLQRRKDR
						VE\PCETQSPGPASGGTLGPSGE
						EAPRPRMSPATIAAKHGEEVSP
						AASDLAMVLT\RGMSL*HQKSS
						RDSLQYSSGYSTQTTTPSCSEDT
						IPSQGRPAGLGPTKPVSIESTYL
		1				QSPQRLKQEWATEGKIQKLPPQ
						IOVVNFQVDSSPSLPLIMVPEH
						MHVS
15003	45371	В	15090	1	777	
15004	45372	A	15091	26	336	KISPMINVFCSESSPSGERRGA*
						TPPSPQSPPCGPKPSS\PPSCSPSR
						NYFLLFQGLCAEHPALAAPAAS
		1				VGTCPSPASGTHAAAPSGPWRG
						RCVHSGPGGRLW
15005	45373	Ć	15092	46	423	
15006	45374	Α	15093	1773	2219	ELDYLYCSNSSAQCPGALLPSL
		1				ALPGPAGSARQAGSPARRLCAP
		1				VQPSGRWPAPGGPGFPGSSCPA
		1				SPTRQ*LLTSGSGCPRPCHHPSG
		1				PPQ*QSTQAWDWVILGPGAHG
		1				PSGQRCERCLPSRPSGPQTHSG
						VGSEPRAVRGPGVSFV
15007	45375	В	15094	1	7916	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15008	45376	Α	15095	16	545	GLPWRQHVRVEECGRS/PLRPC
ļ					į	SPGTPSPS*PLPAATRGPATCQS
						LSGLQPEKTT/SPS/CAAVAHER
						PPGPVPQCPAQGTGLIF*EGWT
						QPQQIRQGPHPVAHE\CAHDLQ
						QAAPGPGGAHQG/PSQPG*PDR
ŀ						QPDAERAVVHPGAQ*DAEIQPG
						LPGEGGPLATRTERSPAQRPSP
						QARG
15009	45377	Α	15096	320	1167	MDLKTAVFNPARDGKLRLLTK
						*\MASKSKEEVSSLISEKTNGAT
						PLLMAARYGHLDMVEFLLEQC
Ì						SASIEVGGSVNFDGETIEGAPPL
						WAASAAGHLKVVQSLLNHGAS
		1				VNNTTLTNSTPLRAACFDGHLE
						IVKYLVEHKADLEVSNRHGHT
		l				CLMISCYKGHKEIAQYLLE\KG
						ADVNRKSVKGNTALHDCAESG
						SLDIMKMLLMYCAKMEKDGY
1		1				GMTPLLSASVTGHTNIVDFLTH
		1				HAQTSKTERINALELLGATLFT
	·					QKEISCDESGRTRDLSPRRKPPH
15010	45378	Α	15097	187	623	KLKTQKNNIGPQACNSQLASVS
						TRSPPPTLSVPIIMTDDSQASQL
						LLAAVTIRKLWSRMAPMLALP
						TGQSGQTGAEGQGKEGERGQS
						PCWKPRLDPKLC/EPRQPAQLL*
						ADVCTGP/CGVRGCPADCRLPC
						LSPSSAQQPKEASRSA
15011	45379	Α	15098	789	1041	
15012	45380	Α	15099	707	1373	SAWPSCHCCHLDILSR/PGDDA
		Ì				CSVQILVPGLKGDAGEKGDKG
						APGRPGRVGPTGEKGEKGDSG
		1				DIGPPGPNGEPGLPCECSQLRK
						AIGEMDNQVSQLTSELKFIKNA
1		ļ				VAGVRETESKIYLLVKEEKPYA
						DPPLSWQGRGGTLSMAQDEAV
		ļ				HGLKGAYLGQARLGRGFIGIHR
		1				PWKRRAPLLLF*TLSPLRSF*KG
ł						DGGGNTQKGSRKKRELAGQIW
						GPCIRR
15013	45381	Α	15100	79	420	WRRLFLLGVLFRGYLGCLRSRS
]						VLGRLKVSDMRIFFFACGCGHL
						STLPSWPLRPSLWLRL*EEQELP
						SLSVPSSGRSASRLL*LGWRGF
		<u> </u>				GPSPSPRLTVQFSGGRRAQGPS
15014	45382	A	15101	3	284	
15015	45383	C_	15102	83	112	

SEQ ID NO:	of peptide	Met hod	SEQ ID NO: in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15016	45384	Α	15103	84	303	EEQELPSLSVPSCLWGPCPRAG
						WLHSLHWPVSHSFSATSTFGM
						AIWT*SLASGIKPQSSSKVSSRIQ
						VPPTM
15017	45385	В	15104	112	462	
15018	45386	Α	15105	326	1323	DRKAINLQVKLEGAVAL*KP\G
		ŀ				EAQLEFGKALGLFQ\STLGHHF
						VNNKEKILGDF\QEATLQAATK
						LTRGRSLVMENMEQLLSMRME
		ĺ				DQSQRNMPLSVVLIREKAHSPF
						ENRKQEQGEGASPENFRAR\QG
						*FLAFKAAPPA*AASGPAAQER
ĺ						SVQEPSVNAPG*IPEGGLHTLA
						GVHHE*ALAWCSPLNKPRFLW
						KRLP/SRTFHFPVEDESAPGSKP
						SGDLLTLLLGGNAAGDFKLK\P
						CLVYPSENPCFLKGSFKPNLPL
						VWCSHKKAWVQPG*QSETPFQ
						KKKKKKKKAWVTMSLFQEWF
						LHFCSTVE/ARCTQYDLPYKGL
						PILDSTPGHPGC
15019	45387	Α	15106	3	375	HASAHASGLQQSYQAVKEKSS
		1				EALEFMKRDLTEFTQVVQHDT
						ACTIAATASVVKEKLATEGSSG
		l l				ATEKMKKGLSDFLGVISDTFAP
						SPDKT\STAMSSPYDGTKARLY
						SLQSDPATYCNEPDGPP
15020	45388	Α	15107	184	377	YCTNNQKVN*TKEIITKRKTPSR
						QKKDSAFSPSHPLTEGLKW\ND
		<u> </u>				LTLSVSLTHDGFVWRRGKGV
15021	45389	Α	15108	3	816	SWEDVGWWRSWLQQSYQAVK
						EKSSEALEFMKRDLTEFTQVVQ
						HDTACTIAATASVVKEKLAIAA
						CSRGACFLCPFSIQTEGSSGATE
						KMKKGLSDFLGVISDTFAPSPD
						KTIDCDVITLMGTPSGPAEPYD
	ļ					GTKARLYSLQSDPATYCNEPDG
						PPELFDAWLSQFCLEEKKGEISE
						LLVGSPSIRALYTKMVPAAVSH
						SE/TFWHR\YFYKVHQLEQEQA
					•	RRDALKQRA/VSGDSCT*SGVL
						TNE*KGSWVRDTERVRSLPLQT
						FSEGVGWREC
15022	45390	A	15109	1	281	

SEQ ID	SEQ ID NO:	1	SEQ ID NO:	I .	li de la companya de la companya de la companya de la companya de la companya de la companya de la companya de	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	detection, \=possible nucleotide insertion)
				sequence		
15023	45391	Ā	15110	1	1194	MGFLHLGQAGLEFLTSGSSTDI
						SEDWEKDFDLDMTEEEVQMAL
						SKVDASGEPFCFWKGYPGGEL
						ETLIRLLQKLFFLKTDPAFVLSP
						STLVKLHLWGNEGLQESLESLV
1						LHDAALVILVPNLEKKGEISELL
		İ				VGSPSIRALYTKMVPAAVSHSE
						FWHRYFYKVHQLEQEQARR\A
						RLKQRAEQSISEEPGWEEEEEE
						LMGISPISPKEAKVPVAKISTFPE
				:		GRTLAPQSPCEENLVTSVEPPA
			,			EVTPSESSESIPKSPLPTRALGTD
15024	45392	Α	15111	213	1031	AGNKPDPWAGRNRTAVLPDVS
	}					VFHREDVGWWRSWLQQSYQA
						VKEKSSEALEFMKRDLTEFTQV
						VQHDTVCTIAATASVVKEKLA
						TEGSSGATEKMKKGLSDFLGVI
						SDTFAPSPDKTI*\RDVITLMGTP
						SGTAEPYDGTKARLYSLQADP
						ATLLY*P\DGPPELFDAWLSQFC
				1		LEEKKGEISELLVGSPSIRALYT
						KMVPAAVSHSEFWHRYFYKVH
						QLEQEQAP\RDALKQRAEQSIFE
						EPGWEEEEEELMGILPIFPKEAK
						FPLAKISYIP
15025	45393	С	15112	414	557	
15026	45394	Α	15113	98	467	
15027	45395	Α	15114	1	155	
15028	45396	Α	15115	16	94	RCPIASEASWTITD*RCPIASEAS
				_		WTITDALGNSYSGGMAF
15029	45397	Α	15116	95	115	IIRHLCND*TPREGCLPSP*PAW
		<u> </u>				SDTFETWVNNQASLQ
15030	45398	Α	15117	3	282	LKPVITCLLQLGLLKPINSPYNS
						PTLPVQKPDRPYSKPNSLP*LNP
						SLLQRNYTSIFLLTLNMPSISCT
		_				TMLLYGQKEISSLHKGPPSLMP
15031	45399	Α	15118	548	1401	SCTPCWHLTGRLWGITAVRQK
						RHVLSVDPKLRHQSRTGKAAFP
						WCLIIAGTPL*LYTHVSRVSDH
						AGMPALVLHPLQQVPLFWGRG
						NTL*KD*SLLSLAC/VQHGLLKP
						INSPYNSPILPVVKPDKPYKLVQ
				1	1	NLRLINQIVLPIHPVGEAPVPLE
]						TGGKSGKDCILWFECHLSHNTI
						EHQVSSSTFRIVVATGVLVSLH
						PQLYMAQNRERHAECLGESKG
						REQESLSGNPENSRFSLLTSRRY
						LYKSTRTIVLLGLAGPLKQKKL
	<u> </u>	L	<u> </u>	L	l	RSQHPSLYKYLQSFRNKDSYR

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15032	45400	A	15119	144	298	WQVPLFWQPFPEPLELWPKA/H LTDSFPDLLGLAAED*RCPIASE APCTQVI
15033	45401	A	15120	449	667	
15034	45402	A	15121	2	111	
15035	45403	A	15122	323	401	LLLGLAAED*RCWIASEAL*TIT DAK
15036	45404	A	15123	2	593	CTLPSH*NLITLTPLNANIPSHG ML*KD*SLLSLACYSTAF*GL*T PLTIPPFYLFKNQTRLTG*FRICP LSTKLFCLSTPWCQTHILSYPQY LPPQPIFLFWISNMLSLLFLCTL YPSLSSLSLGLTLTPIRLSKLPGP YCRKPSQTAPTSNHPSSFSGSW YSVETSYPLSSSIFRKGRTDSWS LKDTPHQAQPPT
15037	45405	Α	15124	354	630	
15038	45406	A	15125	29	354	RFLVSLMLPPAVVMLNHCYSLI FTNALVKRLFAWNMLSELATC AGNLATGPRNARSPGFLLSCVP SVRDPTGNQTVQLTWQPLPEPL ELWPKA/HLTDSFPDLLGLAAE
15039	45407	С	15126	1	576	
15040	45408	A	15127	377	624	RDQLQCLQLWDAGACYTCWK SATGPRNAHSTVFLLSHVPSVW DPTENRTVHLTWQPLPEPLELW PKA/HLTDSFPDLLGLAAED
15041	45409	С	15128	1	462	
15042	45410	A	15129	44	340	PCQTTQGRLLTAGTPL*SFTHVS RVSDHAGTPALVLHP*RQVPLF WGRGNHISGTQELPNT*TAV/V QAF/LPEPPPTG/CLLHVPEIWPL GQGMPAGQDSS
15043	45411	A	15130	242	522	RLVWPTATQRAWSQSEAAAAK TVDLALSARNSATGPRNACSPG FLLSCAPSVWDFTGNRTVQLTR QPLPEPLELWPKA/HLTDSFPDL LGLVTED
15044	45412	В	15131	1	1080	
15045	45413	Α	15132	1	642	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15046	45414	Α	15133	245	1033	TALLLTQSLFGGLFTQTRMKFG
15010	13 11 1	1	13133			AVTRIGGPPLGDQSPVLLFFVPR
						ERSTYDLGPQTDQPKKHLTNFK
						STSFVFSSCIPPP*PSSISLLPPW/T
						TDHAPLTISL/TT*SPLPCSMPIS/
						ASHSML*KD*SLLSLACYSMAF
						*SL*TLLTIPPFYLS*NQRRFTG*
						FRICALSTKLFCLSTPWCQTHIL
						SYPQYLPPQSIILFWISNVLSLLF
						LCTLHPSLSSLSLGLTLTPFRLS
						KLPGLYCRKASQTAPITSVKPK
						FLPYLLPISA
15047	45415	В	15134	146	556	
15048	45416	Α	15135	1	555	LTSFLHCCLSHHYHGAQIHPRH
						FLHLICRCHCHQYCHHAYLDPC
						PQDSHHLHWCWQPLPEPLELRP
						KA/HLTDSFPDLLGLAAED*HCP
		1				IASEAALPPCFPRPDMITSRLQL
		1				QRRRQTALGVRYRNATYRTRT
						ASTNSLRPRGSPFPLALVTNDR
	-					QVRSQLALWKRDRLLQLSPINR
	İ					AISESSFT
15049	45417	В	15136	1	555	
15050	45418	В	15137	499	769	
15051	45419	Α	15138	752	1111	MQPTIGNSQSVSRPTA*MPLAE
1						SHPITQLVGTVTTPRVTPLSIRH
						PWRCRSANAGAMILTMKQLAI
		1				VMRVYQHQVRRSMRPGRNGA
		1				RSGMSAVGAYYSSCIRVMTAL
						QSCTGDDEAVLL
15052	45420	C	15139	1	632	
15053	45421	Α	15140	1	375	PGVRGAQGGPSIPRQCEESAIGP
		1				KFAWIISSFHWKQGMFNMEAP
		1				SSLFFVNMCAVKKQTTWGRPD
		1				RGFIWQPLPEPLELWPKA/HLTD
		1				SFPDLLGLAAED*HCPIASEAPA
		1				LVYTAGLHCFSKPSQ

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15054	45422	A	15141		1340	CSEYEDSSPAPVPATDLSSTLSS SVPQPQDTGTSQQLHPLDPWHE LLRAQELQGATNHKGYSHAEH EHAGLGVQGGNGALAFSNSGH RHAVPTISSGTGRRRTPSSSAFG LLNLHQWFVSGFQAFSDRLKA ALSASLLLRFGDSDWLPSSSAC KCLMLGLHFVIVGNICATLKEK YSSMLHLDVTMKKNGEKRTRL QKRKKGMPPHPAYEDLNIAAIT LPANVVLHQPSGFRTSGQLDPV WWSLDTDAHEIWCQDPGLGSG DFPWEITPLSSYSLLHEKDPPTT SGPQTDQPKKHLTNFKSKTKET GFIHGPKTPAPVTDWEGSLPLV FNH\SGTPL*LFTHVSRVSDHAG MPALVLHP*RQVPLFWRRGKI* LTSPSRCTIIEKSCNSLPPL*DKP QPHLQHTRTSKCLNRSGQAF/L PEPPPTG/CLLHVPEIWPLGQGM PAARDSS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15055	45423	Α	15142	2	2107	IDMIFTPGPPSTPKHKKSQKGSA
						FTFPSQQSPRNEPYVARPSTSEI
		İ				EDQSMMGKFVKVERQVQDMG
						KKLDFLVDMHMQHMERLQVQ
						VTEYYPTKGTSSPAEAEKKEDN
						RYSDLKTIICNYSETGPPEPPYSF
		1				HQVTIDKVSPYGFFAHDPVNLP
		1				RGGPSSGKVQATPPSSATTYVE
						RPTVLPILTLLDSRVSCHSQADL
						QGPYSDRISPRQRRSITRDSDTP
						LSLMSVNHEELERSPSGFSISQD
						RDDYVFGPNGGSSWMREKRYL
						AEGETDTDTDPFTPSGSMPLSST
						GDGISDSTVPLFLSSEILQKQVG
				}		QSITSMLGFLSRGPSMKLCMGL
						ACVLSLWNTVSGIKGEAKKEK
						GMTFLPTTDSKKFFSLLSVTSYS
						SFAFHKFSVAVYNISNLKTVDP
				1		AKFPTRYCYCLNNRTNDLSDFT
						ALLVDIIGNSTSYLTEIFKSTSIL
						SVNQSNESDCIFICVMTGKSGR
						NLSDFWEIEEKYPIINYTFTSGL
		1				SGVLALLLTQSLFGGLFTRTRM
		ĺ				KFGAVTRIGGPPLGNQSPSSCSL
		1				LHEKDPPTTSGPQTDQPKKHLT
						NFKSAARPTFLGQGQVPLNPFS
İ						FTLS/EQVLLS*AARTPQSLISTP
İ		l				QPLISVPQSLISVPQPLLYFSGG
į						QEPPPPPLLCVSSLFSRLASFTM
						GAFTHGTQTPSPTKATAPRYPQ
İ						TGDLSAEWPFTAGEEPVLVPRP
15056	45424	В	15143	1	930	
15057	45425	В	15144	401	1093	
15058	45426	С	15145	1	843	
15059	45427	Α	15146	1	498	
15060	45428	Α	15147	112	669	SLQGRLSDYTPTFQGCQTTQGC
Í						LPWSFTLCSKSRFSGEGETRFKR
ĺ	1					IKACYHSPATAWPFKAYKLSLQ
ĺ					,	FPHFTCRKTRQALQVSSESAPY
İ						QPNCFAYPPRGAEPIHSPILNTS
į	1	ŀ				LHNPLFCSGSQTCFLY/SFLCTL
1	}	ŀ				HPSLSSLSLGLTLTPIRLSKLPGL
ł						YCLKSSQTAPITSVKPKFLPHLL
						PISA
15061	45429	Α	15148	957	1100	

SEQ ID NO:	of peptide		SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15062	45430	Α	15149	303	2028	LGSGDLPWEINPLSSCSLLREKD
						PPTTSGPQTHQPKEHLTNFKSD
1						KGDMFYPWTQN/YQCRSRTGK
						AAFPWCLITAGTPL*LYTHVSR
						VSDHAGMPALVLHP*RQVPLF
						WGRGKLLLARPS*VPILPQPLLL
		l				HPIIFLSPPLLTPGPAYSFVL*LA
						LPHLPSNLLLKRWLEPKA*SSM/
						TLKGLKPVITRLLQHGLLKPTN
						SPYNSPILPVLKPDKPYKLVQD
						LRLINQIVLPIHPVLGIIGLTSSV
						RRDAGQDLKRDRAEFLLGDEV
						HHPHRRRIAEARRLLLGQHFDP
	1					LHRLIGQVLELGEARYAPPVEQ
						HHRLAPARRTGQRLHPLEQFG
						QAGRAQRRNRLGIEHRDRLDRP
						DDGAGNALAGDGDFRRGCLFA
						GIGIRPRYCRYRQSQDDRRPSH
						AAPRPRALPIPHRPATPIADVM
						VMFSVAIMSQLRLVLRHIDGIS
		•				RWPPAVVFITDVKIWLVYPSDD
						CRIRSNDRDDMQGEAPAMSMN
						AAARVGDPIGHSFSQGLFGEAL
						DGLFFARRSEVDMRAGNLGRLI
						ARGLSGGRWTPADGQLTLGSR
15062	45 42 1	<u> </u>	15150	653	1076	DVFINGPPATMTIRSTGQCRQH RSKYPNLVSLCPSPLFPRPDLLS
15063	45431	Α	15150	033	1076	LWPNPLFLHPNLLYLCAPIPYFH
		l				APTSYLCTPTPYFHDPTPFPLFW
						KELATSARNLATRPRNACSPGF
	:					LLSCVPSVRDPTGNRTVQLTW
						QPLPEPLELWPKA/HLTDSFPDL
						LGLAAED*CCPIASESP*TSDILG
						RDTLLLALKVQTVVLQTACGE
						GHVAGNCGRPLETEGSLQLTAT
						KKLRDSVLQPKSPEFCQQFTRA
						WNRTQVPDETEAPAGTYAAQS
						GDLPWEINPLSSCSLLHEKDPPT
						ASGPQTDQPKEHLTNFKSGFRG
1						VRPRRDACLGPSPLAASPAFLG
						KGQVPQPLLSVSLPLLRLSGGQ
						ETPNPFSFTLSGKSAFLEEQVPQ
						PRISVPQSLISTPRPLISVAQSLIS
						APQPLISLCPNPLFPCPDLLSLHP
	!					NPLFPRPNPFPAFLEGACYKCQ
						KSGHQAKECLQPGIPPKLRPICA
						GPHWKSDCPTHLAATPRAPGT
						LAQGSLTDSFPDLLGLAAED
15064	45432	C	15151	234	449	

SEQ ID NO:	SEQ ID NO:	Met hod	SEQ ID NO:	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15065	45433	A	15152	2095	2663	LGLGDLPWEINPPSSCSLLREKD
ł				1		PPTTSGPQTYQPKKHLTNFKSA
ļ	ļ					CFERIKACYHSPAKAWPFKAYK
				İ		LSLQFPHFTCLKTRHAI/PS*FSD
						LCLINQIVLPIHPMVPNPYTLLC
						SIPPYTMHYSVLQLIRAFFTIPLH
						P*\PSLY*VSLGLTLTPIRLSKLP
						GLHCRKASQTTPITSVKPKFHP
						HLLPIST
15066	45434	Α	15153	678	1189	LGSGDLPWEINPLSSCSLLCEKH
						PPTTSGPQTDQPKKHLTNFKSE
						TKETCFIHEPKTLAPVTD*GRQP
						SLGV*SLQGRPL*SFSHISRVSD
						HAGMPALVLHP*RQVPLFWGR
1						GKYPNPFSPCLLPLLCFSGDRG
						KYPNPFSPCLYSFSAFLGTGQVP
1.50.45		↓			1.400	QPLLSLSLPLLCF
15067	45435	Α	15154	649	1428	QCISELQFLLASTVRQTPATSPA
						HKNFQTPELQRPGVPPEPPPTG
						ACYTCRKSGHWAKECPQPGIPP
						KA/HLTDSFPDLLSLAA\ED*HC
						PIASEAPETITDAELPVTLTVEA
		1				HLHPGEINSHVAHTKPVWWSL
						HRDAHEIWCCDSDRGTSLRRPI PCPPALCSVTKIHLRPQVLRPTS
						PRNISPISNPAAPHQAELGPNPS/
						SSLCSSTL*SFYHLPSSHLVWLT
						VSFRD
15068	45436	Α	15155	240	1208	LRSGDLPWEINPLSSCSLLREKD
				}		PPTTSGPQTHQPKEHLTNFKSC
	İ					KQ/DLFLLSSPTSLTIPQPLSPFN
				İ		LGATLQSLPSLNFNSFHFLVETK
						ETRFIRGPKTPAPVT/VLGRQPS
						LGV*SLQRCLSDYTPTFQGCQT
						TQGRLPWSFTLSGK/CPLFWGR
						GNTL*KD*SLLSLACYSMGLLN
						PINSPFPSSHFTCPKTRQAVTS*
						VQDLRLINHIVLLPIHPGGAQPH
						TLFCPQYRPVVLPIILFVYVKHA
						FFTIPLQPLIPSLSLPSLRLTLTPI
						RLNKLPGLCCCKAFRDSPHYLN
						QAQISSSSVTYLGIILIKAHVLSL
1.5055	15.10=	 	1.5155	1110	1.622	PIVSDW
15069	45437	Α	15156	1174	1623	KFGLVQLTLGKPLPEPLELWPK
						A/HLTDSFPDLLGLAAED*HCPI
						ASEAPYTI\TDAELRVTLTVEGK
						SVPFLINTEATHSTLPSFQGPLSL
						ASITVVGIDGQASKPLKTPQLW
						CQLRQYSFKHSFLVIPTCPVPLL
	l		<u></u>	L	L	G*DTLTKLSASLTIP

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15070	45438	L	15157	1 1	1610	 MGLGKSLCILKDAVLALREIKG
13070	43438	Α	13137	1	1010	ISFQYAPNTNRLWGYRQAKQT
						QHLSGISTLDCPQRYNGDDTGS VWALEPSCMLWACPLVFPSEIY
						YFISEDVWKHLDAQEKFAAVM
						GPSRRTSVRAVWKGNVGLEPP
						HRVPTGALPRVAVKRGPLSSRP
						QNGRSTYGLHCLKDTPTQEDW
						LVSVLPEGSRVGVDPLIIPTALG
		İ				HGRTTFLKVLDFQGPAPCPAA
						MGVRLMAGPQCVSSDYWKKM
						AKVLRSAGHHLIPVKENLVDKI
						WTDRPERPCKPLLTLGLDYTGI
						SW\KDKVAD\LRLKM\AERNVM
						W\FV\VTALDEIAWLFN\LRGSD
						VEHNPVFFSYAIIRD*ETFMLF\I
	ŀ					DGD\RIDAPSV\KEHLLLDLGLE
ŀ						AEYRIQVHPYKSILSELKALCA
						DLSPREKVWVSDKASYA/V*SE
						TIPKDHRCCMPYTPICIAKA\VK
						NSA\ESEGMRISDEKSAGNLTED
						PFYVIDRTLITTGVVCNSTEKSS
						LMSVASADTAENLWHLFFHILP
						SERALTRKLRVLKLGPVNYSVQ
						NPTAQLHQQGPHLGVA
15071	45439	A	15158	1	756	męznągomee m
15072	45440	A	15159	1	1677	
15073	45441	Α	15160	1220	2620	GPTEPHTGELWLESGGCPSGTK
						LPEEGTGSNLCCSTASAGDTQA
						NRVCSGPPTNHSRPAADRPVRR
						KINKQKTATSTSTKRTSTQRPH
						PKVTNDHSSSPAREQNWMEN\E
						FDELTEVGFRRWVITNSSELKK
						HVLTQCKEAKNLEKRHSLGTK
						EPAALKGRTQYWQDSSPADLR
						PWLCATLLKQYKPTVFSHSTVA
				1		VHPRAGFLNLSTVDQHPVLDEP
						IAEAVCIIADTNKWSVQVATSQ
			ļ			RKVPHNLKLGQDVLVSSQVSSL
						LQSILQLYKLHLPADFVGDILLD
						NNLKCVFQLGSILPVTFRSHDP
						YDGEGTDELRTAACSPDTEQLA
						RQSGTYPVSLGQATVLEDSKA
						ALPGLGPTLMAVTVVTTEAAV
			1			DMVVGDRGEERGGTLVAPGTR
						GLTEEIWFPVPSPSRWWSSHPV
						VAFITLAKYGPTSSPPVFHHGPN
						IIRHGAQGAASAGDSGSSSRCD
						RLPCRQNA
	I .	1	L	L	L	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15074	45442	Α	15161	666	1536	FSSAGDTQANRVCSVPPTNHSR
						PAADRPVRRKVNKQKTATSTS
						TKRTSTQRPHPKVTNDHSSSPA
		ŀ				REQNWMEN\EFDELTEVGFRR
						WVITNSSELKKHVLTQCKEAK
				:		NLEKRVRVNCLT\RIT*P*RRTI\
		ŀ				KTLMELKNTARE\LHEAYTSINS
		İ]		WINQAEERLSEIEDQLNEIKRED
						KITEKMKNSTTVRVAASMQSK
						LLQGVAEEGPLRLTRSASFSAEI
					:	SMCHQWLFSTFGNGKSRNRYH
						CEKAHIEGNQLVEETGRAFKSR
						HRYGTGPLHKFEIHEYTELHHV
		l				LMLQGKC
15075	45443	С	15162	104	364	EMEQUICE
15076	45444	A	15163	162	467	
15077	45445	В	15164	1	1122	- · · · · · · · · · · · · · · · · · · ·
15078	45446	A	15165	188	456	
15079	45447	Ā	15166	485	1580	GEGYKADLAAATVECPICQQQ
	15	` `	15100	100		RLTLSPQYSHIPQGDQPTTW*Q
						VDCIGPLLKGQRFVLTGIDTYS
ŀ						GYVFAYPVCNASAKTIISGLTE
		l				CLIHCHGIPHSIASDQGTHFMA
		1	ļ	i		KEVWQWAHVHGIH*SYHVAD
ŀ		1				YPEAAGLIEGWNGLLKSQLQC
		1		}		QLGDNTLQGWGKVLQKPMYA
		l		;		LNQHLIYGTVSPIATIHGSRNQD
		1				WNGSSLRGWTFTILFTAPRREP
		ļ				GTYIASDQRTHFMAKEVWQW
]				AHAHGIHWSYHVADYPEAAGL
]				IEGWNGLLKSQLQCQLGDNTL
		ŀ				QGWGKVLQKPMYALNQHLIY
		ł				GTVSPIATIHGSRNQGMEVEVA
į		}				PLTITPGDPQAKFLLPVPMTLCS
		1				AGLEVLVPEGGTLPPGHTTIPLN
15080	45448	В	15167	10	2352	TODE VENTE OF THE OFFICE OF THE OFFI
15081	45449	A	15168	2	497	WKPIAMRKSRRRQEKLLPVQK
	13 1.17	١,,	10100	[EGQPKEGEREEGGPVQAQKVP
						NRLHLHWTT/HLPSFFQLPLLLD
		i				SLWRALKFVVLEAIVKMIRFNG
						VLGKCEAIRAQALHEGGPPLPL
						AVFHNRRLQIVACSRISARLRGS
						PPPKSNNEQSSLSTDLRTDSQRL
						PPLSRNAQQKG
15092	45450	D	15160	1	1911	FFLSNIAQVAO
15082	45450	В	15169]1	1711	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		·
15083	45451	Α	15170	1	1569	MPIGKVGEYYIKGTLRATKESE
						QQSSALELSSDRVYLNEKEPGD
						QPWNAAFVFRGCGVALALRPP
		1				HLGWGEPRWAGFRGAFGFPFS
						TLNPSNWSLTLIIQGCKQEGFYS
						PSMRPRVVPKRGSAPKVRAEEN
		t				AALPSRCPPGPLPVAQPRGLLM
						GAHVSSAPAVGVGAASCHRLC
ļ						PGFPDRPRPTLRSCIMNAVVGG
						CDLRGQVYDPTGGWANEGTR
						DLSAVQGVPNHPGGRPCPTPQS
						SRQLPVVSRRPVVRPGGEAPPP
				•		LDILPAALALHQSLLTPQDPRE
						DRPSPADLWAQDSWVGQPQEL
						PVGSRVRTGTEPDTHSGSVCTS
						RQGKDGSGPLARESSVRVILQG
						PSSAWMMFPEYTPGPRARAGT
		1				AVVGAMARQTVNVCLTKVRL
						QHPEGIPGGGRGPLVPKPVCTC/
ŀ						IVDVKGSRV\WHVHICMSMGC
						RHVCLCGRSVGSWRMCTAGQT
						PKLARQASIELPSMAASSTKSW
						WETGEVQAQSAAKTPSCKDIV
						AGDMSKKSLWEQKGGSKTSST
						IKMDSVTT
15084	45452	Α	15171	328	641	LLMRSWSLPSAPRGFSCSCLQP
						QYPCRTESWPQTEPVPSPVSHR
						LLVLATAMLGSSME/RLAG*LG
				ŀ		GPASGLDGLGVLFQPLINLGDG
		ļ				QVWLGFLHTVTLLYGAV
15085	45453	В	15172	120	1458	
15086	45454	Α	15173	1	440	IQQDSQAPLYS\IKETQRANTYL
						VEWEPGAETAFKTLKQALVQA
						PALSLPTGQNFSLYITERAGIAF
						GVLTQTRGTTPQPVVYLKDGEP
						IEHDYQQIIAQTYATQDDLLEV
						PLANPDLNLYTDGSSFVVNGIR
		<u> </u>				RAGYAIVSNVTVLE
15087	45455	Α	15174	288	376	NHWLLLTMDPRIQQDSQAPLY
		L				S\IKETQRA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
İ	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15088	45456	A	15175	777	1548	SKIQRDSQAPLYS\IEETQRANT
						HLVEGEPEAEIAFKTLKQALVQ
i						APALSLPTGQNFFLYITERAGIA
						LGILTQTRETTPQPVAYLSKEV
						DVVAKVWLHCLRDGEPIKHDC
						QQIIVQIYAARDDLLEVPLANP
l	,					DLNLYIDGSSFVENGIRRAGYAI
						VSDVTVLENSLAAVTLQNRQG
						LDLLTAEKGGLCTFSGKECCFY
				1		TNQSGISSPLEDTTTAGPFLHPI
						QQEVARAIIGQFPTAFGVSCLE
		1				GRLRGEASWTSWVE
15089	45457	A	15176	1	1359	MGPEPKNTHPGSCTCPLACSSS
						HKGFECLAVEQGSYTPVASPIV
	ŀ	ł				GVKENQEILDLLELKYNQTRSA
	ł	1				VTRCVLGAASTARIKQAEEGGR
		ŀ				SRLAESSGLRLPFVLELPALEHQ
		ı				TPSSSGFELLDLHQRLATWSQA
						FNYRLKDTLLASLLLRFWDSD
						WLPCSSVCSWPIVGLHLVIIERN
						YSYQQGTETVVSSWATGVSAA
		1				ASVSATIPINRVQGHCGFLGKR
						CFYCCIGECNHSDHQGPGTVAG
		1			ļ	SWARGVSATASQIQVDLGKFS
			<u> </u>		1	DDPDRYIDVLHLTWRDVMLLL
	İ	1				DQTLAFNENNVALAAAQEFGD
		ł				TWYHRPVNDRMTAEQRDKFPT
		ŀ				DIKTAAGIPWNHWLLLTMDPQ
		ŀ				TQRDSQAPLYS\I*ETQRANTYL
		1				VEWEPEAE/TSFKTLKQALVQA
						PALSILTGQNFSLYVTERARLAL
		1				GVLTQTCGTTPLYQDELRKKTL
						RHRDSEGSGFNQLEASAD
15090	45458	Α	15177	368	5192	AAPAPQLGLGPCKLASHLLSQS
						HLSQSVRSEAEQKAGASPASSI
						QREPLFKDQLKAPLPEQKRRSL
						LTVPLLTFRAQKCGPGSPLVAR
		:				GGATDSWLVSEEFRLSDSTAPP
		ļ				TELRGQEATGITKPPASLHPQPH
						AWDWARGRDPRRHLLGTPELQ
						EHSEKTTTETLPADMGKLRTW
						ERKEATAGQAAEGAAGGIHFE
						ATSLLRPSSYVPQTQGALSTTY
					1	KEAKMDRDLHLTPILAAQRAR
						GPRIHSFPSDEMIEPS
L		L		<u>L</u>		OLKIHOLLODEMIES

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
15091	45459	Α	15178	76	969	KKSTALVIDNGSCMCKAGFAG
					:	DDAPQAMFPSTVGFPGHQGVM
						VDMGQKDSYVGNEAQSKRSIL
						TLKYPIKHGIITNWDNMEKIWH
						HFYNELHGAPE*QPVLLTEDPL
					1	NPKDNREKMTQIMFETFNTPA
						MYVAIRDVLSLYAFGRTTGIVM
						DCGHRVTHTVPIYEGYSLPHAI
						LHLDLTGWDLTEYPCILLFSEK
ļ						SYELPDGQVITIGNEWFWCPEV
						LFQPSFPCMESCRIHQTTINSIKK
•						CDVDIRKD\NTVLSGGTTMYPG
						IADRMQKEITALAPSTMKIKIIA PPERKFSVWIGCS
15092	45460	A	15179	1	451	FFERRISV WIGCS
15093	45461	A	15180	130	227	LMKMKKVM*RA*KEEERRTPP
,	10.01					AEDDFEKDLEWLINENEKSDAS
		l				IIEVYIANSIHNNYITKCFAIL
15094	45462	A	15181	104	1715	KEEERRTPPAISVIPAAAFYTAY
						FLHKQKMNENKDTDSKKSEEY
						EDDFEKDLEWLINENEKSDASII
						EMACEKEENINQDLKENETVM
			:			EHTKRHSDPDKSLQDEVSPRRN
		1	į			DIISVPGIQPLDPISDSDSENSFQ
					1	ESKLESQKDLEEEEDEEVRRYI
		l		İ		MEKIVQANKLLQNQEPVNDKR
		ļ				ERLLKFKDQLVDLEVPPLEDTT
		1				TSKNYFENERNMFGKLSQLCIS
						NDFGQEDVLLSLTNGSCEENKD
						RTILVERDGKFELLNLQDIASQ GFLPPINNANSTENDPQQLLPRS
						SNSSVSGTKKEDSTAKIHAVTH
						SSTGEPLAYIAQPPLNRKTCPSS
						AVNSDRSKGNGKSNHRTQSAH
						ISPVTSTYCLSPRQKELQKQLEE
						KREKLKREEERRKIEEEKEKKR
						ENDIVFKAWLQKKREQVLEMR
						RIQRAKEIEDMNSRENRDPQQA
						FRLWLKKKH\EEQMKERQTEEL
						RKQEECLFFLKGTEGRERAFKQ
						WLRRKRMEKMAEQQAVRERT
						RQLRLEAKRSKQLQHHLYMSE
1						AKPFRFTDHYN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15095	45463	Α	15182	1881	2040	MKAGWRQTQHLHDAGQLLHL
	1.5 1.5					LPGEQGVARVQLRRDAAQAPH
						VNGHVVGVAQDHFWGSVEPA
						LDVRVYCGGKTGGKQRERREG
						RCGVEILMVGERKMNENKETD
						SKESEEYKDESEKDLEGLTNEN
	:					EKSDASIIEMACEKEENINQDLK
						ENETVIEHSKQLSDPDKSLQDE
		1				VSPRRNDFISVPSIQPLDPISDSD
				:		SENSFQESKLESQKDLEEEEDEE
						VLPRSSNSCVSGIKKEDSAAKID
		1				VVTHSSTGEPLAYI/ASTYKLAY
		1				IPPLNRKTCPSSAAISDRSKGNG
						KSN/HGQSAHISSVTSTYCLSP*
						QKEL/QKREQKREKLKREEEQQ
						KIEEEEEKKRENGIVFKAWLQK
						KREQVLEMRRIQRAKEIEDMNS
						RVSKTSLKNKFIQI
15096	45464	Α	15183	873	4292	MVVSLGRVSSLSEDSDPTGRVG
	ŀ					R*VDCKVDRSIWAGVWISEDAP
	ļ					SEA*EIGGLGRK*MLSGGPQSPS
	ļ					KHVHIL*DQDSSSIPEPMGPCCP\
						PHAPGKEPGTSCSIDTDGKLSG
						VSGDDS*T/GTDGGESEEILLPQ
						*QLPVTLPSLPLPG*VRYTSRSR
						SRSRSTRDPGRKSRPISGLRGLC
				İ		GDGATS/PLTPRHAPARDVQ/PG
						VSMAFAGDR\GDGGRRGGASQ/
						PAPPGPAVPRPCAQYSARAMPA
						SGRRES
15097	45465	В	15184	1	2214	
15098	45466	Α	15185	3	361	
15099	45467	Α	15186	1	1035	MKIHKSGGLKATPQLYASQFSG
						VAILATWGPHRPRNSSTSFSTQS
						KLRRPDESSVDFQKLMLCKEAT
						KKSKEKEPGMAFPQGHLIFSDV
		1				AIKFSVEEWKCLNPEQRALYRE
						MMLENYRNLES/VG*LFKIHDG
						VLINRARQYRSVPHRDIGKT*FL
						LPRNQERYSGL*VSVARN*KK
						WP*STHDRNQRVN*VHTGEKP
						QKCNECGKTFSQKSYLQCHHR
						LHTGQKPYK*E*CDKVYSCRSQ
						LKTHRRIHTG\EKAYKCKVCEK
		1				AFWDNSCLSCQKRVHIGEKPYT
						ALVMHKAIHTGEKPYTCNECG
						KAFSRKANLALYHRLHTGEKP
	-	_	1.5.0-	<u> </u>	10.45	YKCEECDKVYSRRSHLERHK
15100	45468	В	15187	1	1347	

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15101	45469	A	15188	<u> </u>	1446	MWRETKRYAAVLLLTPHYSAS
13101	73702		15.00			LLPDW/PNLYLHQSGHHVPGVY
						LKFG\SSNPIYPCVDSAKTVFPN
		1				APLHDRVQSVRRMGPFKGVFR
						KVFSNLGFEYSLSTWATMGSPG
						SLGVSPFGKTLFVESASGSLDA
		l	1			LWPTVEKEISSHKNYQKHSEKQ
		1				IYDVRIELTGLNALPNISLNFYK
						NSVSKLLNRKKGLIHLLQPFILR
						LCVSLHKNREENLGITIQDVAM
						RLSALPNIVCRLYKKTASEVLK
						EWSTRWDEYTHHKEVSQKALP
						TLSVKIYKKNICQLLNQNNGPN
						SEMNALITKKSVKSASGYLESI
						AALVEKEISSYKNYTEAFYNLL
						CDVYSHVTEMKLSSIEQIGNTL
						FVESASGVWRALRPRGKEVNDI
						FIDQIANSVLVDSSKDICKRFEA
						CGAMVKKENLHIKTRQKISEKL
						HWPFGAPRLTSLTGPEKTGKGE
						HPGAKARGELSGAPPLIHTRKL
	ŀ					SLMSNVETLLLEKLQGYTGAH
15102	45470	Α	15189	1	966	BEINGIN BIEBEBREQUITOINI
15103	45471	A	15190	157	531	GDHTTFLCLLMVNAEFKYNFK
		-				TIIS*DLLKKDSGQYGHFTIFFRS
	ļ					MSIGCFSIYLCIFNFFHQCFAVF
						LVEIFTFLVKCIPQVDSIEDTTFG
						HPEEATPHPFSFFMRLEQFSHIF
						SSWPVSFTAH
15104	45472	A	15191	9	494	PTDRPRPRALAGARGPPGRAPR
			1			VPPPAGGPRPSQYGHGGRPNPR
						HGWHDAFCPNNTPYR\ERPLDS
						RYLTPGPLDMRGDSGFRTGRGS
						DVGSLKPTDTRRNGKIYLNGS/
						NVFYYQQRLHPVHRGDGARRG
						FSGRTCLHRMVC*YEQTGHQS
			ļ			DQT*KARSAYG
15105	45473	Α	15192	1361	2038	
15106	45474	Α	15193	1376	1906	LAFPDGMASLIMMSLPAMRTL
						AWAMAVSGNCIPEGQFSALSC
						ADFGISPKLIL*SSSAPN\YLAMG
						ISCPRADDLPEAGTAETDRACR
						SIPQYGTDLGWYPATAAGPQSE
						SVQGAV*PHRYEAG*ST/VWKP
						LKKK*KPLPRR*LARWQMIRFR
						KPSPLTTSLKLTCAWRFLDFRW
						LE
L	L	ь	<u> </u>	L	L	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15107	45475	Α	15194	11	1388	MIEHPRRRDNYRQSGPPVADKP
						TTKGEPLISDIVSDAQQANLLIP
						VDETPPVINDEQSTSTPLTTAQT
						MALAAVADKNTTKDEKADDL
						NEDVTASLSALFAMLPGFDNTP
						KVTDAPSTVLPTEKPTLFTKLTS
						EQLTTAQPDDAPGTPAQPLTPL
						VAEAQSKAEVISTPSPPSISGVT
						RSKMRQREWLVLPPEDSLVIRR
						SNKTSQLPAFNVPLSAAKAAAS
						KPWRASNTACRNSRPSKVCSAE
						GEQWFSASGRGEIHSQSPARDM
						AQMCKLAAEGHFAEARVINQR
						LMPLHNKLFVEPNPIPVKWACK
						ELGLVATDTLRLPMTPITDSGR
Ì						ETVSGDETYLEAAPLAELHAPA
						GMILPVTSGDYAIPVTNGSGAV
						GSYSAITEPGAGSDVGSLKTTY
						TRRNGKIYLNGS/NVFYYQQRL
						HPVHRGDGARRGFSGQTCLHR
						MVC*YEQTGHQSDQT*KARSA
						YG
15108	45476	C	15195	1	1905	
15109	45477	Α	15196	285	1706	YARVHSGRSSIRRSIPVSICIRSSI
						RGCVPVVHCIRSSIRRSIPVDHL
						YQIIHTKKHTRCPFVSDHPYEE
				1		APDYSASLLPDWLPVPSSKVVI
						TYPGFTSICGYYTTSSKACFSVF
		ļ				AWCFVTSDVPGRFSDEDEDDSE
						MWSLTALMGVMGGWKTSARS
						EVSILCSTSSKVIHSSYPYADRA
						FQVWSLDARFAHINKPFGVDRF
						SGEAPSRAITTMYGVKVLNRPA
						TGTAHRLVPPDVPVHPHGGESN
		ŀ				KPRVQTTVSSGISRLSIPISASAL
						QNVPDKTHDISNITDTVYHCPL
						SQSSRWYWRFVPVRSPPLSNGQ
		1				TAVRPGGSRAGAAGGPLSEYS
						GSGFGVMKWTKDQLTVTVKV
						HHLPEVVEFLYYKQGGWLSVL
						FGNDERKLNGHYAVYYVLSRE
						KGTKCWI/TDSMDYRQR/PAPT
	1					NDAETYEFINELGARKTTVVPI
						GPLHVTS/DEPGHLRLFVDGENI
						IEANNE*IHRTQC
15110	45478 45479	A B	15197 15198	3	1396 1399	
15111				11		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15112	45480	Α	15199	283	796	FVGVKSAGNRLKTIDELANESR
						SSVGLTQAALKTLESSGAIRIER
İ						RGRNGSYLVEMDNKALLTHVD
				1		INNVVCAMPLPYTRLYEGLASG
		ŀ				LKAQFDGIPFYYAHMRGADIRV
						ECLLNGVYDMAVVSRLAAESY
						LPQKGLCLAL*LGPPPYVGE\YQ
						L*QVKAKSEYVKRVGHV
15113	45481	A	15200	48	3376	2 (
15114	45482	A	15201	49	681	DPGADRRGSVWGLSDLRNTYN
						ALYLTHSGRLSECPTPANEEKE
						PTNHNRTLTNQTQKEKKKTRT
						KDCTTRNKNSTRPGSGIIAGPSY
		l				YTVAGPPSAIDPRWLRPTPPAL
İ		l				DPQ*MPHTSKRGKRADQSQPN
		İ				ANKPNTEREKKDKNEGLHHKE
						QEQ\QRPGSGIIAGPSYYTVAGP
						PSAIDPRWLRPTPPALDPQTEPL
						IFQQLEIDHYVAGLTHEVMSLS
						HLTLVMRWGSVSAAHLAHWA
	ĺ					RCRWISTELTPNSITSTSSPLKD
						CSSSPAMEQSWLENDFDELTED
ł						KKTCLCDH
15115	45483	Α	15202	3	587	
15116	45484	A	15203	3	774	MVLMAGFTAGNEKGELVVLG
						RNGSDYSAAVLAACLRADCCEI
						WTDVDGVYTCDPRQVPDARLL
		l				KSMSYQEAMELSYFGAKVLHP
						RTITPIAQFQIPCLIKNTGNPQAP
ŀ						GTLIGASRDEDELPVKGISNLN
						NMAMFSVSGPGMKGMVGMA
						ARVFAAMSRARISVVLITQSSSE
						YSISFCVPQSDCVRAERAMREE
						FYLELEESLLEPLAVT\ERLPII\S
						VGSDGNPPPCGGISPKFF\AALA
			1			PRPISKFVALCSGIPF

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15117	45485	A	15204	1474	1511	HS*VPANLYMELKACFTGA/NE KGELVVLGHNLAVSYFIALAAS LRADCCEIWTDVDGVYTCDPR QVPDARLLKSMSYQEAMELSY FGAKVLHPRTITPIAQFQIPCLIK NTGNPQAPGTLIGASRDEDELP VKGISNLNNMAMFSVSGPGMK GMVGMAARVFAAMSRARISV GLITQSSSEYSISFCVPQSDCVR AERAMQEEFYL\ELKEGLLEPL AVTERLAIISVVGDGMRTLRGIS AKFFAALARANINIVAIAQGSSE RSISVVVNNDDATTGVRVTHQ MLFNTDQVIEVFVIGVGGVGG ALLEQLKRQQSWLKNKHIDLR VCGVANSKALLTNVHGLNLEN WQEELAQAKEPFNLGRLIRLVK EYHLLNPVIVDCTSSQAVADQY ADFLREGFHVVTPNKKANTSS MDYYHQLRYAAEKSRRKFLYE HNTFVSPHIRPLIERRGRSSTRD KWCARPSTERQQDNSAARILSC QQGNTSFEHTSSIRVGTTNSLN
15118	45486	Α	15205	198	462	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15119	45487	Α	15206	31	1981	RWLMRGMGIALPAAMASTVSP
						STIAETPEPPPLSDW/LRASG/DV
						WVHLLGLDPHLSSLLLSICLFSR
						SLIWIF*LQP*SV/ALAFQMGAS
						LFASNIGVIEGEGPQAKPL*CLT
						VSYFEMLNNLNAQGPKGKLSP
						CQCLSLQVMTMPEYLKKRWTL
						GQRSQWPQMHVSFSS*LFPF*A
						DIFAGAIFIKQPSGEGKYMRTH
						G*NALVLFFLSLGGLEVVIINIG
						CLRFQ*DISFYSDAFLAFNEVGG
						YESFTEKYVNATPSVVEGDNLT
						ISASCYTPRADSFHIFRDAVTGK
						GR*LPLKSSPGLFHECVTFSRQV
						IVQRCL\CGKDMSHVKAACIMF
						LFV*WLLVLPIWASLDSLLNSSP
						DMVACVVPSECVISTLSGLARQ
						CPATNQCLYPFPAPGLRGLMLS
						VMLASLMSSLTSIFNSASTRTK
						AQWK*L*CFCLMFKLCCFLLRI
						FVLLLTVVSIVWVPLVQVSQNG
						QLIHYTESI***LT*ISFSYFLNILI
						FFLLLLQGAFWGLMVGLAMGL
						IRMITEFAYGTGSCLAPSNCPKII
						CGVHYLYFSIKVPFYYTIANICL
						LLMHFCMVPVQLYRLCWVLR
						NTTLTRISSSFVI*IFCNHLLIFVP
						DYPEKSRGCLKKAYDLFCGLQ
						KGPKLTKEEEEALSKKLTDTSE
15120	45488	В	15207	1	1392	
15121	45489	A	15208	57	1523	
15122	45490	Α	15209	473	589	LPRQLALSCVVHCQTLVPAIRCI
						DWGWLF*W*SVTPVM

SEQ ID			SEQ ID NO:	I .		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 109/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		07/340,217	sequence	or pepuae sequence	deterior, a possible macretified insertion,
15123	45491	Α	15210	1	1297	MGKIMSSYQHVDLIVDCDFEA
						YHRQPSSGTKSVVTGPSAQALI
						TEKIVTYHWDQHVGDVLVLNI
						YCVINVLDHTVKAVIDDRFMG
1		1				GITTINLQLMAIIVRYFFASGSA
						YIVAMLPVFAMLANVSGAPLM
						LTALALLFSNSYGGMVTHYGG
		l				AAGPVIFGVGYNDIKSLCFDER
					:	TRIDVADTYASNWRSRCNQTIF
		ĺ			1	NGECPNTRQHIAAVRRGIDTLL
						MNDYLRKQIIDISTRIAGTANN
						GDFTGQWITAANTVNFQIVAGT
						HNCHQYFIALRLIFGILKYIQDR
			:			AREGLKQRDLAEALSTSPQTVN
						NWIKRDALSREAAQQISEKFGY
						SLDWLLNGEGSPKKDLESNIPP
						ESEWGTVDAWDKNTPLPDDEV
						EVPFLKDIEFACGDGRVHDEDH
1			ŀ			NGFKLRFSKV\VQPRVGC*QG*
		ļ.,				G*LLYLDSV**GDPLEST
15124	45492	Α	15211	62	301	SWMSSPRAFSSSTCLETAAPTR
						MATALTSAACRLPPAAGSGSP
						WAPAPYRWRARRRTGPMSDR
15105	45493	A	15212	1	1665	Q*RTSQPCPEPAAASR MQDSAHLRRLPPAASRRVWLA
15125	43493	A	13212		1003	LGSCPLPLESSPAHRAYERPSVA
						HQPAVPGARRSLEVPSEVPLTV
						PALTPPRIPAMLTAVCGSLGSQ
ł						HTEAPHASPPRLDLQPLQTYQG
						HTSPEAGDYPSPLQPGELQSLPL
						GPEVDFSQGYELPGASSRVTCE
						DLESDSPLAPGPFSKLLQPDMS
						HHYESWFRPTHPGAEDGSWWD
						LHPGTSWMDLPHTQGALTSPG
						HPGALQAGLGGYVGDHQLCAP
1						PPHPHAHHLLPAAGGOHLLGPP
					İ	DGAKALEVAAPESQGLDSSLD
						GAARPKGSRRSVPRSSGQTVCR
						CPNCLEAERLGAPCGPDGGKK
						KHLHNCHIPGCGKAYAKTSHL
						KAHLRWHSGDRPFVCNWLFCG
						KRFTRSDELQRHLQTHTGTKKF
1						PCAVCSRVFMRSDHLAKHMKT
1						HEGAKEEAAGAASGEGKAGGA
						VEPPGGKGKREAEGSMASSSPD
						SPCSCDCFVSVPPASAIPAVIFA
						HELGPTPGRGAGGGVCPRRHS
						HSWEPAPGL*AVGGGGGASGQ
			1			AGDRGCHA*EAGRS/SCSRRGS
:						GL*RAGTS*ISASS
——				<u> </u>	L	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15126	45494	Α	15213	3	471	GRPSSPGVHGSSTQTPAHVGSF
						PGCGPIPPD/ACRRSKIRGPGGW
						GGVGG/CRDGRNSR/DLRNLPC
						PFLFLPGP*AAAPSPPWPAAGA
						APCGQCLDGWPHSYSHGHSSA
						AGRGGICRSPPLLAGELRPPGIQ
			•		·	YNGRTEVRGILSGSPCPPFSGAG
15127	45495	Α	15214	3	449	PHQDLEEAMVLL*PPSP/PAWPT
]						MRTRKRPSSKVSSTSRPLRKSIM
						TTYAVPSRAPTPA*PSASKPTNA
						FS/HMVGSEPRSHAHLDGRHRD
						SR*RKPRPLSGPRTLGGKSLRGS
						AGAGRGGVPRPRPLPENLDAPL
						WNSPGNSAGAAVQLA
15128	45496	Α	15215	26	1126	EGPSSEAAHPGGTPREQDGAM
						DPARKAGAQAMIWTAGWLLL
						LLLRGGAQALECYSCVQKADD
]						GCSPNKMKTVKCAPGVDVCT\
Ì						EAVVAVETIHGQFSLAVR\GCG
						SGLPGKNDRGLDLHGLLAFIQL
						QQCAQDRCNAKLNLTSRALDP
						AGNESAYPPNGVECYSCVGLSR
						EACQGTSPPVVSCYNASDHVY
				•		KGCFDGNVTLTAANVTVSLPV
						RGCVQDEFCTRDGVTGPGFTLS
						GSCCQGSRCNSDLRNKTYFSPR
		ļ				IPPLVRLPPPEPTTVASTTSVTTS
						TSAPVRPTSTTKPMPAPTSQTPR
						QGVEHEASRDEEPRLTGGAAG
						HQDRSNSGQYPAKGGPQQPHN
		ļ				KGCVAPTAGLAALLLAVAAGV
15129	45497	A	15216	1401	1983	AQRAQALECYSCVQKAD/DGC
						SPNKMKTVKCAPG/VDVCTEA
						VGAVE/SHPFWIWPRPFNAAPPP
						LSAWPGPYDTALQARPLY*QL
						QARGAES*ELVH/GQFSLAVRG
						CGSGLPGK/NDRGLDLHGLLAF
						IQLQ/QCAQDRCNAKLNLTS/RG
						LDPAGTSPPVVSCYNASDHVY/
						KGCFDGNVTLTAEPTTVASTTS
		<u> </u>				VTTSTSPQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15130	45498	IA	15217	1	964	MNVKGDTQGNTRVRVDNIGG
13130	13470	ļ'`	13211	•	501	VGAQTVNGIELIEVGGNSAGNF
						ALTTGTVEAGAYVYTLAKGKG
						NDEKN\SPSVYRP/EAGSYISNIA
						AANSLFSHRLHDRHSCLQPLKI
						LVTGGAGFIGSAVVRHIINNTQ
					<u> </u>	DSVVNVDKLTYAGNRESLADV
						SDSERYVFEHADICDAPAMARI
						FAQHQPDAVMHLAAESHVDRS
						ITGPAAFIETNIVGTYVLLEAAR
						NYWSALDSDKKNSFRFHHISTD
						EVYGDLPHPDEVNNTEELPLFT
						ETTAYAPSSPYSASKASSDHLV
						RAWKRTYGLPTIVTNCSNNYG
						PYHFPEKLIPLVILNALE
15131	45499	В	15218	44	405	
15132	45500	Α	15222	1	498	MTRSLLSVFEEDAGTLTDYTNQ
						LLQAMQRVYGAQLNLLHTELA
						KQL\ETQWFYLSYNSEKRISQTL
						SAMWRGRAAQA/TQGHWVWG
						MVEAVCSKVSTLKDLFGLASN
1			ł			EHDLSMAKYSRLPKKKENEKL
			:			AAFGQPTWCDPAERVGFLEPV
						GPDPGAKSAPSPRDRASTL
15133	45501	Α	15223	1	1224	
15134	45502	Α	15224	1	1353	
15135	45503	Α	15225	1	2580	
15136	45504	A	15226	1	3117 1260	
15137 15138	45505 45506	A	15227 15228	1	2142	
15139	45507	A A	15229	1	1413	
15140	45508	A	15230	1	4049	_
15141	45509	A	15231	1	381	MLMLNIGFKNFQFEHLKCSHGS
15141	45507	``	13231	1	301	YSEDIGFPNAGALSGAMFLLEE
						KHLLFGLKEEERLRASIRRESQ
						QRRMREKQHQRGLSASY\WNLI
						DTMRRRKARSP/YSLAAIKNRY
		:				KGGIRDQKPSYDSLNLCMT
15142	45510	Α	15232	1	467	DHSSSPAREQNWMEN\EFDELT
						EVGFRRWVITNSSELKKHVLTQ
						CKEAKNLEKRLGELLTRITSLE
						KNINDLMELKNTARELHEAYTS
						INSWINQAEERLSEIEDQLNEIK
	[REDKITEKMKNSTTVRVAASM
					<u> </u>	QSKLLQGVAEEGPLRLTRSASF
15143	45511	Α	15233	1	1140	
15144	45512	Α	15234	1	2640	
15145	45513	Α	15235	1	714	
15146	45514	В	15236	1	546	
15147	45515	В	15237	1	735	
15148	45516	В	15238	50	1603	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		·
15140	45517	<u> </u>	15239	1374	2220	OA ENGLININIZZOVIJET DEEC
15149	45517	Α	13239	1374	2220	QAENSHININKKDVHSETPSEG
						HQRKRPKDHSSSPAREQNWME
		İ				N\EFDELTEVGFRRWVITNSSEL
						KKHVLTQCKEAKNLEKRLGEL
						LTRITSLEKNINDLMELKNTAR
						ELHEAYTSINSWINQAEERLSEI
						EDQLNEIKREDKITEKMKNSTT
						VLVSTWGSQLEQNNLPHIISPLF
]	İ				HSTHDLNISFYGGCLLQACSSK
						LLQGLAEGGPLRLTRSASFFAA
		1				KLCLTKHFLPTEMILIKSRKTTP
1				•		DTQPRRLYLLQGVVDQTSRKG
Ĺ						PTALNIRQKTFAAAPPHR
15150	45518	В	15240	1	1311	
15151	45519	В	15241	46	212	
15152	45520	Α	15242	3	371	EPLPPEPPAVPSEVELQHLRKEL
	İ	ł				ERVAGELQSQVKNNQHISLLNR
]					RQEERIREQEERLRKQEERLQE
		1				QHEKLRQLAKPQSVELK\SQEV
		1				QSLQQQPDHF\LGHL\QQYV\AT
						LSAARWAAYQQLD
15153	45521	Α	15243	392	558	
15154	45522	В	15244	52	689	
15155	45523	В	15245	400	603	
15156	45524	Α	15246	5	1158	RKFWRWMVVMAAQHSQAAQ
						GQRR**VSKGP**PGPMGGPVL
						GHWWHSGGISLAVPSLPPPGKS
						SVFLFLFSASFSDMKLNPGTVSP
						RLIFGSYDDGFLVETVVKFDFQ
						LWQKTPLHRAAGERRRAEQRG
						KLSFVMP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
E	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15157	45525	Α	15247	1	1808	MAEETQHNKLAAAKKKSATGF
13137	43323		13247	1	1000	HREGPTSSATLKDLESPCQERA
						VVLDSTSVKISRLKNTIKSLKQQ
						KKQVEHQLEEVTAPDNLVPWV
		ļ				GLSWGIGGILGACLLLCHLCLP
						LEKKANNERQKAERELEVQIQT
						LIIQKEELNTDLYHMERSLRYFE
						EESKDLAVRLQHSLQCKGELES
						ALSAVIATEKKKANQLSSRSKA
						RTEWKLEQSMREEALLKVQLT
						QLKESFQQVQLERDEYSEHLKG
						ERARWQQRMRKMSQEICTLKK
						EKQQDMRRVEELERSLSKLKN
1						QMAEPLPPEPPAVPSEVELQHL
						RKELERVAGELQSQVKNNQHIS
						LLNRRQEERIREQEERLRKQEE
						RLQEQHEKLRQLAKPHCAFSRS
		İ				*TMRTRAHLQLEQQVKELQE/Q
						ALARVFLQVELKSQEAQSLQQ
						QPDHYLGHPAAEHLEAASQQN
						QQLTAQLSLMALPGEGHGGEH
						LDSEGEEAPRPMPSVPEDLESR
						EAMVAFFKSAGASAQEKQAQL
						QEQVKEQRVCCQRLAHPVASA
						QKEPEAARGPGAPGPGGESSSF
						MDHLKEKADLSE/PGEKRTLLH
						P/PTG/GDRRHQKTHHLLSEPGG
						CAK/DAALGPGHHQAGAQGGD
15158	45526	A	15248	2085	2700	YPGKRGLEWT*ANSNRPAAEG
15156	173320	``	13240	2003	2700	PDE\SSSLPATEQSWMENDFDE
						LRKEGFRRSVITNFSKLKEDVO
		1				THRKEAKNLEKGLDEWLTRINS
						IEKTLNDLMEMKTMA*ELHDT
						CTSFSSRVDQVEERLSVI*DQM
						NEMK*EEKFREK\RVKRNEQSL
						QEIWNYVKRPNLHLIGVSEIDR
						ENGTKLENTLQDIFQENFTYLA
						•
	L					RQANIQIQEI

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15159	45527	Α	15249	378	1744	QTERNSININKKDIHTKTPSVRH
1	1					HHQRPKLHKTTKMGKKHSRKA
						ENSKNQSGSLPPKECSSSPAVE
						QSWTENEFDEFTEEGFR\RIESQ
						MNEIKGEEKFREKRVKRNEQSL
						QEIWDYVKRPDLRLIGVPDSDG
						ENGTRLENTLQDIIRENFPNLAR
						QASIQIQEIQRTPQRYSSRRATP
						RHIIVRFTKVEMKEKMLRAARE
ŀ						KELKIKKLTQNCLTTWKLNNLL
						LNDYWVHNEMKGEIKMFFETN
						ENIDTTYQNLWHTFKAVCRGK
						FIALNAHKRKQERSKIDTLASQ
						LKEIEKQEQTHSKASRRQEITKI
ŀ						RVELKEIETQKSLQKINESRSWF
						FEKINKIDRPLPRLIKKKREKNQ
						IDTIKNDKGDITTNPTEIQTISRE
İ						YYKHLYANKLENLEEMDKFLD
						TYTLSGLNKEEVESLNTSITGSE
						VEAIINSIPTKKKSRTRWIHSRIL
						PEVQGGAA
15160	45528	Α	15250	935	1269	LIEGKLTERNSININKKDIHTKT
						PSVG/PPSSKTKDRSSLPAIEQS
						WMENDFDELTEVGFRRSVITNL
•						SKLKEDVRIPHKEATNLEKRLD
						EWLTRINSIEKTLDDLMELKTM
15161	45529	Α	15251	401	497	KTFSPFYFS*RRRTVRDINGFTE
						Y*EPEKLLG
15162	45530	В	15252	220	940	
15163	45531	Α	15253	1	1476	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15164	45532	Α	15254	1	2423	MNGKNKTNIYQEPTVCQALHW
						TLARIGKEEPRRGQRGEGLLCS
		i				RERQPRGLSSPGTAAPAAPAAP
						TPPALPGHRTRSGSPEGPEPILR
						RGGSADAPGRVKTWVNRPAAF
						VGKCLGDGSADTACDQAMKG
						QRGSNRKRKTSHSLTFPTSSDM
		İ				HAGGLSAAHPCAHSRAPPPASP
						SPYKGSLGTRTDGLTDSRADSA
			l.			ASGARALGVDGQPQPSPPRRAP
						PRPVGADGSSRGPQPGRRRVER
						RGPATQPGDSRALPEPRGVPAV
		İ				HPAGSGSEWERPPPAAPSPEHR
					· ·	DKMLPGLRRLLQASARRPQPFP
						EAARWVWSPCFSSGTLCPSFQS
						ERPLLPFQGASFPGTTLHQRLCP
						VAALHGSGKGQKWQQSHCYA
						AAVMEEPPAPGRSSESLQGPLS
						WAGGAPASACLLLMLLALPLA
						APSCPMLCTCYSSPPTVSCQAN
		l				NFSSVPLSLPPSTQRLFLQNNLI
						RTLGARHLWVQPAHPVALLQQ
		1				PLHHLPGHFPPL\QALEELDLGD
						NRHLRSLEPDTFQGLERLQSLH
-						LYRCQLSSLPGNIFRGLVSLQYL
		l				YLQENSLLHLQDDLFADLANLE
	İ					PTSFLHGNRLRLLTEHVFRGLG
						SLDRLLLHGNRLQGVHRAAFR
						GLSRLTILYLFNNSLASLPGEAL
		l				ADLPSLEFLRLNANPWACDCR
	1					ARPLWAWFQRARVSSSDVTCA
			:			TPPERQGRDLRALREADFQACP
15165	45533	В	15255	47	482	
15166	45534	Α	15256	1	636	MRDPNTKRSRGFGFVTCATVE
1						EVDAAMNARPCKVVGRTVEPK
						RAVSREDSQRPEDTEELHLRDY
			1			FEQYGKIEVIEIMTDQGSGKKR
}						GFAFVTFDNHDSMDKTVIQKY
						HTVNGHNCEARKALSKQEMAR
						ASSSQRGRSGSGNFGGGRGGGF
		1				GGNDNFGRGGNFSGHGGFGGS
1						HGGGGYGG/SGDGYNGFGNDG
l						GGGSYNDFVNYNNQSSHFGPM
L			I		·	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
1,0.	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15167	45535	Α	15257	1	890	MSKSESPKEPEQLRKLFIGGLSF
						ETTDESLRSHFEQWGTLTDCVV
						MRDPNTKRSRGFGFVTYATVE
						EVDAAMNARPHKVDGRVVEP
						KRAVSREDSQRPDYFEQYGKIE
						VIEIMTDRGSGKKRGFAFVTFD
						DHDSVDKTVIQKYHTVNGHNC
						EVRKALSKQEMASASSSQRGRS
						GSGNFGGGRGGGFGGNDNFGR
						GGNFSGRGGFGGSHGGGGYGG
						SGDGYNGFGNDGSNFGGGGSY
						NDFGNYNNQSSNFGPMKGGNF
		ŀ				GGRSSGPYGGGGQYFAKPRNQ/
						GGYGGSSSSSSYGSGRRF
15168	45536	Α	15258	1	286	
15169	45537	Α	15259	1	1674	
15170	45538	Α	15260	235	1370	KREGEGRLADTFVSLLLSARGR
						RRRSIVKVSLHPAVMSKSESPK
						EP\EQLRKLFIGGLSFETTDESLR
						SHFEQWGTLTD\CVVMRDPNT
						KRSRG\FGFVTYATVEEVDAAM
]				NARPHKVD\GRVV\EPKRAVSR
						EDSQRPGAH\LTVKK\IFVGWH*
ŀ		İ				RKTTFWGGKKKRAKKHHLRD
						YFEQYGKI\EVIEIH*LDR\GQWP
						RKRGFAFVTF\DDP\DSVDKIVI
						QKYHTVNGHNCEV\RKALSKQ
						E\MASASSS\QRG\RSGSGNFGG
						GRGGGF\GGN\DNFG\RGGNFS\
		1				G\RGGFGGT\RGGGGYGG/SVG
		İ				DGYNGFGNDGSNFGGGG\SYN
						DFGNYNNQ\SSNFGPMKG\GNF
		l				G\GRSSGPYG\GGGQYF\AKPR\N
		i				QGGYGGSSSSSSYGSGRRF
15171	45539	В	15261	1	660	
15172	45540	Α	15262	2	486	KMRQRERAQGKK/VAPAPAVV
						K/KQEAKKVVNPLFEKRPKNFG
						IGQDIQPKRDLTRFVKWPRYIR
						LQRQRAILYKRLKVPPAINQFT
				}		QALDRQTATQLLKLAHKYRP/G
						DKGALAKLVEAIRTNYNDRYD
						EIRRHWGGNVLGPKSVARIAKL
						EKAKAKELATKLG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15173	45541	Α	15263	3	845	PPKMPKGKKAKGKKVAPAPAV
						VKKQEAKKVVNPLFEKRPKNF
						GIGQDIQPKRDLTRFVK\WPRYI
ŀ						RLQR\QRAILYKRLKVPPAINQF
						TQALDRQTATQLLK/LVAHKYR
			}			P\ETKQEKKQRL\LARA\EKKAA
						WQKGTSPTKRPP\VLR\AGS*HP
İ					:	VTTLVE\NKKASAGWWIAHRR
			}			GFPSSLVVFLP\ALCREKWGSPY
						\CIIKGKARLG\RLVTRKTCTTV
						AF\TQVNLGRQRALLLKLVEAI
						RTNYNDRYDEIRRHWGGNVLG
						PKSVARIAKLEKAKAK\ELATK
15174	45542	Α	15264	1	5796	
15175	45543	Α	15265	1	2212	MPRVEKNIPGRRNSKGFEDPLE
						VTQDVTREWAKKVVWKREKA
						SKINGAYFCEGRVRGEAIRIRT
İ						MKMRQQATLTMTVDKGDNVN
ŀ						ISFKKVLIKEEDAVIYKNGSFIH
						SVPRHEVPDILEVHLPHAQPQD
						AGVYSARYIGGNLFTSAFTRLI
		l.				VRILPPQPENIKISNITHSSAVIS
						WTILDGYSISSITIRYKVQGKNE
						DQHVDVKIKNATITQYQLKGLE
						PETAYQVDIFAENNIGSSNPAFS
						HELVTLPESQAPADLGGGKML
						LIAILGSAGMTCLTVLLAFLIILQ
		1				LKRANVQRRMAQAFQNVREEP
						AVQFNSGTLALNRKVKNNPDP
						TIYPVLDWNDIKFQDVIGEGNF
				•		GQVLKARIKKDGLRMDAAIKR
						MKEYASKDDHRDFAGELEVLC
						KLGHHPNIINLLGACEHRERGD
						RRHLENQVTPTLILRFSNTLSKR
						HTRRLYPEPGSEGPTPTEPRSLL
						AQQSEIKLHGGSEPPLLIFRQTG
						SGVDLQQTPTNLQLRVLTVRR
						KTDKRKGHPHQKPICSSPSSKT
				1		KEGFRRSNYSKLKEEVQTHGK
						EVKNLEKKLDKWLTRITNAEK
						SLKDLMELKTKARELCDECASL
				1		TSQFDQLDKRVSVMEDQMNE
			1			MKQEEKFREKRIKRNEQSLQEI
						WDYVKRPNLCLIGAPESDGEN
1						GTKLENTLQDIIQENFPNITRQA
		1				NIQIQEIQRTPQRYSSRRATPRHI

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15176	145544	 	15266	<u> </u> 1	743	DDL LEDOTCSCANDLOOTDTM O
15176	45544	Α	15266	1	/43	PPLLIFRQTGSGVDLQQTPTNLQ
						LRVLTVRRKTDKRKGHPHQKPI
						CSSPSSKTKEGFRRSNYSKLKEE
	}					VQTHGKEVKNLEKKLDKWLTR
						ITNAEKSLKDLMELKTKARELC
						DECASLTSQFDQLDKRVSVME
						DQMNEMKQEEKFREKRIKRNE
						QSLQEIWDYVKRPNLCLIGAPE
						SDGENGTKLENTLQDIIQENFPN
						ITRQANIQIQEIQRTPQRYSSRR
			-			ATPRHIIVRFTKVEMKEK\MLR
						AAREK
15177	45545	Α	15267	1	618	
15178	45546	Α	15268	200	618	EVSRNNTRTKTSYKQDMNTSK
						GF*KDTST*PVT/KERSSSPATEQ
						CWTENDFDELREEGFRRSNYSE
						LKEEVRTNGKEVKNLEKKLDE
ļ						WLTRITNAEKSLKDLMELKTM
						ARELCDERTSLSSRCDQLEERV
						SVMEDEMNEM
15179	45547	Α	15269	158	594	SGSSICCSPISTVLQPPLLTPRQT
ŀ		1				GSGATSSK/SPTDLQLRECSSSP
						AKEQSWMENDFDELREGFRRS
						NYSELKEEVQTHGKEVENLEK
		1				KIDEWLTRITNAEKSLKDLMEL
						KTMAREQCDECTSLSSRRDQLE
	<u> </u>					ERVSVIEDQINEMK
15180	45548	Α	15270	3	1453	LSDLELAAQSIIFIFAGYETTSSV
						LSFTLYELATHPDVQQKLQKEI
						DAVLPNKERSSSPATEQSWTEN
						DFDELREEGFRRSNYPELKEEV
						RTHGKEVRNLEKKIRQM\VTRII
1						NAEKSLKDLMELKTMARELRD
				1		ECTSLSSQFNQLEERVSVMEDQ
						MNEMKKEEKFRRKRIKRKEQS
						LQEIWDYVKRPNLRLIGVPESD
						GENRTKLENTLQDIIQENFPNLA
						RQANIQIQEIQRTPQRYSSRRAT
						PRHIIVRFTKVEMKEKMLRAAR
						EKEIQTTIREYYKHLYANKLEN
						LEEMDKFLDTYTLPRLNQEEVE
		1				TLNRPITGSEIEAIINSLPTKKSP
						GPDGFTAEFYQRYKEDLVPFFL
					1	KLFQSIEKEGILPNSFYEASIILIP
						KPGRDTTKKENFRPISLMNSDT
						KILNKILANRIQQHIKKLIHHDQ
						VGFIPGMQGWFNIRKSINIIQHI
						NRTNDKTHMIISIDAEKAFNKIQ
						QPFMLKTLNKL
15181	45549	С	15271	111	269	
10101	1.55.17		1	I · · ·		<u>L</u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15182	45550	Α	15272	94	4505	YLDMMSFVQKGSWLLLALLHP
			1			TIILAQQEAVEGGCSHLGQSYA
		l				DRDVWKPEPCQICVCDSGSVLC
						DDIICDDQELDCPNPEIPFGECC
		l				AVCPQPPTAPTRPPNGQGPQGP
		1				KGDPGPPGIPGRNGDPGIPGQP
						GSPGSPGPPGICESCPTGPQNYS
		l				PQYDSYDVKSGVAVGGLAGYP
		İ				GPAGPPGPPGPPGTSGHPGSPGS
						PGYQGPPGEPGQAGPSGPPGPP
		ŀ				GAIGPSGPAGKDGESGRPGRPG
						ERGLPGPPGIKG
15183	45551	Α	15273	1628	1832	RRRRKARERRRLAAEQGRRMK
						EQCKDGFSE/QASGALKLMGSN
		ŀ				EGEFKAEGNSKFTYTVLEDGCT
						KHTG
15184	45552	Α	15274	1	362	
15185	45553	Α	15275	1076	1240	LRQLL*LCPSWRLCLLGGWQL
						HLFF*RHHFVHLVSQRCSIAMS
						PSPRRCFSLHL
15186	45554	В	15276	480	1167	
15187	45555	Α	15277	3	379	
15188	45556	В	15278	13	369	
15189	45557	Α	15279	112	1112	WPPASPSASVIRTVKEFALTNPE
						KSSTKETERKETKAEEELDAEV
1						LEVFHPTHEWQALQPGQACPC
						RNPTYGLNLQTGEREAKLQYE
		ŀ				DKFRNNLKGKRLDINTNTYTSQ
						DLKSALAKFKEGAEMESSKED
				}		KARQAEVKRLFRPIEELKKDFD
						ELNVVIETDMQIMVRLINKFNS
						SSSSLEEKIAALFDLEYYVHQM
						DNAQDLLSFGGLQVVINGLNST
						EPLVKEYAAFVLGAAFSSNPKV
[Q/EWEAIEGGALQKLLVILATE
						QPLTAKKKVLFALCSLLRHFPY
[AQRQFLKLGGLQVLRTLGAGE
						GHGGARRARGHTALRPCHREK
	<u> </u>	<u> </u>	1.550:			CSPRRRLS
15190	45558	В	15280	54	385	
15191	45559	В	15281	274	776	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15192	45560	A	15282	362	1323	 HQLGRSALLDTRTVSMQKAKN
						GDA VTEPQVA/ENEANGKDTA
:						EVDLLTKELEDFEMKKATARA
						VTGILASHPCSTDVHIINLSLTF
į						HGQELPSDTKLELNSVRHYGLI
						GLNGTENSMLLSAVGKHGVPIP
						EHIDIYHLTREMPPSDKTPLHSL
						VTVSSPRQRDDMSDGHLCTAR
						RHTDIRDDRSPDAPDTARSLAR
						STITTRRIFALATPSRTLQQVCS
						ALAYTSHPAARTQPLASTALTR
						RYMLPSPQLPSLRDHRRTNPLH
						SAVATIPHTTTGEADVSLLCIGL
						STLHYALVYHVSVLTALSAQS
						MSATDASIIFFEPKR
15193	45561	Α	15283	3	537	
15194	45562	A	15284	1	455	MPRPEKLGKTQVPPLAECLTKD
					}	SFLGQKQASLGPLTFRDVTIEFS
						LEEWQCLDTVQQNLYRDVMLE
						NYRNLVFLVISS/HFTQDFWPD
						QSIKDSF/QRIILRTYARCGHKN
						LRLEKIVKVSMRKHQIIHTGKK
1.510.5	1776	<u> </u>	1.500.5		1007	PYKCEECAKLLASPQPLENMR
15195	45563	Α	15285	1	397	FAVLMAHYDVQEEDPVLTVIT
						YMGLSVSLLCLL\LAALTFLLC
						KAIQNTSTSLHLQLSLCLFLAHL LFLVAIDQTGHKVLCSIIAGTLH
ŀ						YLYLATLTWMLLEALYLFLTA
						RNLTVVNYSSINRFMKKLMFPV
15196	45564	A	15286	191	1332	AIETQAITDNCSEERKTFNLNV
13170	13304	`	13200		1332	QMNSMDIRCSDIIQGDTQGPSVI
						AFISYSSLGNIINATFFEEMDKK
						DOVYLNSOVVSAAIGPKRNVSL
			İ			SKSVTLTFQHVKMTPSTKKVFC
						VYWKSTGAG/CSQWSRDGCFLI
						HVNKSHTMCNCSHLSSFAVLM
						ALTSQEEDPVLTVITYVGLSVSL
ļ						LCLLLAALTFLLCKAIRNTSTSL
						HLQLSLCLFLAHLLFLVGIDRTE
						PKVLCSIIAGALHYLYLAAFTW
						MLLEGVHLFLTARNLTVVNYS
						SINRLMKWIMFPVGYGVPAVT
						VAISAASWPHLYGTADRMLAF
İ						KATAQLFILGCTWCLGLLQVGP
						AAQVMAYLFTIINSLQGFFIFLV
						YCLLSQQAKHKFTPMEVPRQN
						HRCLQQN
15197	45565	A	15287	263	397	GRPPAPAARVSGPASVSAG*KR
						VSASGSVGVRSMRRGSALLPPN
				1		F

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion) MKFHDRGQMQPKPSPAEAVNS
						APQREPEGSILRAADPLVWAAK AMSASR/RGS/CGGSSADPRHTE RTPTEPDADTRFQPAETLAGPK TRAAGAG
15199	45567	Α	15289	5	545	SFMTEAKSSPNPAQQRLSTQRP SESRKVPSSELQTLSCGLQRPCL HPGRPGPDPYYASASTRTRSPS ARHAVWGRRGCYPEA/AVRAR EGPHPTLAPASTYGLASPPRAD SAHRPRHRSLTRLLPNRKLGGS SADPRHTERTPTEPDADTRFQP AETLAGPKTRAAGAGGQPWDS ANASPP
15200	45568	В	15290	121	822	
15201	45569	Α	15291		1308	MLRMVGIEISNGQWSTPPCQVL HLEGNDIYNPKEQGLGNTCRW GWMAAGTRRNLCWAPFPFLNP VASLVLHSTIQIALQESKDRVL GTLINSPLVTSTDKANWSLRLA LEVGDGGGLVGLSPQPVGSGA MSRCPNCTRKLIGSGISPTSSTL AEEQGSVVKTHSHQGGARCSK EKGWQILLILVAGLCQLVVVV QNVGVRKIPRKYLAALRVAIGH ALAIERRRIRVDVADQAQDTDR DDRSFGRDRNRDSDKTDTDWR ARSATDSFDDYPPRRGDDRFGD KYRDRYDSDQYGRGYWYEYR DGPCRDMDRYG\GSGRRAFG SGY/RRDNDYRGGRDRYEDRY NRWDDGSCSSRDDFSRDDCRY NDRRPPQRPKLNLKPHNTPKGD DSSASTSQSSRASFVFGGAKPV DRAARETKVEEWLQKEQDKLQ HQLDEAKLE
15202	45570	Α	15292	2	293	
15203	45571	C	15293	56	298	
15204	45572	Α	15294	1	273	TRGPWCDSVLRGCSLEQRSFIS VRLLSYLSACRHPMEDSMDMD MSPLRPQNYLFG\SLGAGAKDE LHIVEAEAMNYEGSPIKVTLAT LKMS

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15205	45573	I A	15295	3	11024	CGGIHSVLRGCSLEPAFFYLRSA
1.5205	1.0075	-				FSPTISACRHPMEDSMDMDMSP
						LRPQNYLFGCE\LKADKDYHFK
						VDNDENEHQLSLRTVSLGAGA
						KDELHIVEAEAMNY\EGSPIKVT
		l				L\ATLKMS\VQPTVFPWGAFEIT
						PTQWSLRLKCG\SGPVHI\SGQH
		1				FSKLLEEDAE\SEDEEE\EDVKL\
						LSISGK\RSAP\GGGSKVPQKKS
						KTLLLMKDDDDDDEED\DDED
						DG*WMDFGWMEEAEEKAPSE
						RNPIRDTP\AKNAQKSNQNGKD
		l				SKPSSTPR\SKGQESFKKQEKTP
		l				KTPKGPSSVEDIKAKMQASIEK
		1			1	GGSLPKVEAKFINYVKNCFRMT
						DQEAIQDLWQWRKSL
15206	45574	В	15296	1	1107	
15207	45575	A	15297	1	1968	MLASMGVVAKEMKYLSSPYPQ
						HAFHTRIKSLMTVNFQAQLWL
						PNEAGPTGDGDWGGDTHGCPS
						FSPHLPRFIQKQPWGTGRRLLQ
	Ì					EHFSCKSLVPVIPSRDIEAFHLH
	·					ADLILNTLPSLVIIIGQIESHSEEL
						RARVSTYESGEDTIQCITVTWK
						PKYGATRWKDPGSPDGPLEER
						RLEDLSTRNNSIRLFRGCGYRV
						KQNSSQRPNKVCGLPDTGVLL
						QTKCGLVRNSQRSAQTNTKQH EELFSTAGIPENLSRRQEVLPGH
		ŀ				GFTGWVGWTWKLTSGEGVTF
						HHLSATHKLSHASCRQGNFKP
						KGAMSVTRSTVHPHLAFLSAEI
						NGKCGRPCVFLGIWAINCGMAI
						LYERLSALVCVSVSERHLDYT
						MRFLFDIPPSTLEARIGSPLWAL
						SLHPLRSSTVTSSVELILPECTV
					; 	AVLEIQYVFLGDTILFTTVGKA
						RSGTRDRDGGRAQIMQVLQGK
						GRELDAAAQPEGQLLREVRVL
						GVPFIPRARVDAWLVHTVAVG
						SADEAHGLLGAAAASSTGGAG
			[ASVDGGSQAVQGGGGDPRAAR
		1				SGPLDAGEEEKAPAEPTAQVAD
						AGGCASEENEVLREKHEAVDH
						SSQREENEERVSALKENSLQQN
						NDDENKIAEKPDWEAEKTSESR
L _		L				NERHLNGADTSFF\SLEDLFQLL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15208	45576	Α	15298	1	2104	MKHLKRWWSAGGGLLHLTLL
						LSLAGLRVDLDLYLLLPPPTLL
						QDELLFLGGPASSAYALSPFSAS
1						GGWGRAGHLHPKGRELDPAAP
						PEGQLLREVRALGVPFVPRTSV
						DAWLVHSVAAGSADEAHGLL
						GAAAASSTGGAGASVDGGSQA
						VQGGGGDPRAARSGPLDAGEE
						EKAPAEPTAQVPDAGG\CASEE
	İ					NGVLREKHEA\VDHSSQHEENE
						ERVSAQKENSLQQNDDDENKI
						AEKPDWEAEKTTESGNE\RHL\
						NGTDTSFSL\EDLFQLLSSQP\EN
		!				SL\EGISIWGDIPLP\GSISDGHEF
						FKHIYHVNFQPRLIS\QD\VNLH
						EAILLCPNNTFRRDPTARTSQSQ
						EPFLQLNSHTTNPEQTLPGTNLT
						GFLSPVDNHMRNLTSQDLLYD
						LDINIFDEINLMSLATEDNFDPI
				1		DVSQLFDEPDSDSGLSLDSSHN
				1		NTSVIKSNSSHSVCDEGAIGYCT
						DHESSSHHDLEGAVGGYYPEPS
						KLCHLDQSDSDFHGDLTFQHVF
						HNHTYHLQPTAPESTSEPFPWP
		l				GKSQKIRSRYLEDTDRNLSRDE
	ļ					QRAKALHIPFSVDEIVGMPVDS
						FNSMLSRYYLTDLQVSLIRDIRR
						RGKNKVAAQNCRKRKLDIILNL
						EDDVCNLQAKKETLKREQAQC
						NKAINIMKQKLHDLYHDIFSRL
						RDDQGRPVNPNHYALQCTHDG
						SILIVPKELVASGHKKETQKGK
15209	45577	Α	15299	2	342	RRRFRVAAAAVGTA*AVVLSA
				1		RPSETWETRGRR*\RICPSSWP*
						GI*VNNG*K*RRHSVCQCSLWF
						PARCIKPYHSMARTQPGTRNKE
						NGPAGPTALDNVASSDDTGRH
						RPQT

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15210	45578	A	15300	710	1831	IRMKSKEIIARCIKPYHSMARTQ PGTRNKENGPAGPTALDNVASS DDTGRHRPQTTQLAPGFAHPLQ LASFRRMVLFLSGEGRSRGGPE LQFPASCRRGEGSPGVRESGSG GIAATSTPNYPPNQDSKEHDIR GAEHQKQEQPAKPPHTARSAY PPQKSSYPANAKATRHSPETAA AKEARAPAAAQPQRHQPNPSP APHTRPATAATRQPERRVPSPT HRHPAATRLSPRRQSPSPRPHH DRRGFPRLAETLQHPMCPLPLV ASAGHHRHHRLLLLLAPQAPER EASDEI\VFSGRSRSRGCPTEFQE SAMCFPNPGLPDSCGESQAVTI LILRKFQKVIWVIEVPLDYKKG SWEYFSRMETIIMPFENIQSE
15211	45579	A	15301	77	1041	IPSCVVSGIAAAF/HPGLASDAP ARASSWWTHVEMGPPDPILGV TEGFKRDTNSKKMNLGVSAYR DHNGKPYSIHKAEAQIAAKNLD EDYLSIGRLAEFCKASAEVALG ENSEVSKSGRFVTVQTISGTSAL RIRASFLQRFLKFSQDVFLPKPT WGSRTPIFRDTGMQLQGYRYY DRKTCGFDFTDTVEDISKTPEQ SLLLLHACSHNPTGVDPGPKQQ KEIATVVKKT/RNLFALFDMAY QGFASGNGNKDAWAVRHSAN\ IRHCRCQSYTKNMVLHSEGVA GFTMVCKDADEAKRVESHLKI LMCPMYSNPPLNGTRLLLPF
15212	45580	A	15302		1263	MTGRQSLSPVLSLSSLDSMSFT TRSIFSTNYLFLGSVQVSNYGA RPASCMASVYAGAGGSGSRISV SRSTSFRGGMGSGDLAAGMAG GLAGMGGIQNKKETMQSLNDR LASYLDRVRSLETKNRKLESKI REHLEKKGLQVRDWSHYLKTI EDLRVQIFANTVDNAGIVLQID NACLAADDFRVKSLMTNVTWL PLETEMEALKEELLFMKKNHEE EVKGLQAQIASSELTVEIEKSTT VVTTQSTKVGDAEMTLTELRR TVQSLKMDLDSMRNLKASLEN SLREVEARYTLQMEQLNRILLH LESEMAQSRAEGQHQIGEYEAL LNIKVKLEAESATYHRLLDDDD FNLGDALDNSNSMQTI/QKTTT RRKVVSETNDTKVL

SEQ ID	-	1	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15214	45582	A	15304	184	370	
15215	45583	Α	15305	40	1411	RGDPRVRRRTQSLSPVLSLSPDS
						MSFTTRSTFSTNYRTLGSVQAP
						SYGARPVSSAASVYAGAGGSG
						SRISVSRSTSFRGGMGSGGLAT
						GIAGGLAGMGGIQNEKETMQS
		į				LNDRLASYLDRVRSLETENRRL
				ļ		ESKNPEHLEKKGPQVRDWSHY
						FKIIEDLRAQIFANTVEQCPTTV
		İ				L/LRIDSA\RLA\ADDFRVKYE\T
						EAGPCAQSVENNIHGLCKVIDD
						TNVTRLQLETEIEALKEELLFM
						KKNHEEEVKGL*AQIASSELTV
						EVDAPKSQDLAEIMADIRAQYD
						ELARKNREELDKYWSQQIEEST
						TVVTTQSAEVGAAETTLTELRR
		ļ				TVQSLEIDLDSMRNLKASLENS
						LREVEARYALQMEQLNGILLHL
						ESELAQTRAEGQRQAQEYEAL
		1				LNIKVKLEAEIATYRRLLEDGE
						DFNLGDALDSSNSMQTIQKTTT
		<u> </u>		<u></u>		RRIVDGKVVSETNDTKVLRH
15216	45584	Α	15306	392	1449	GPWEMLMTIKSHPDAERQPQR
ļ		l				VKKLTQVFWSSASLRER/GKTY
		Į.				LFWPKLKKAAEIPAAYREALDE
						ALGISGGGRDVGERVYKGPSM
						MANPQKPVLEALSSNKYLHILC
						AKHCDRHEGYKDEIKVKEASS
						RVNGCVEAMLSAPLPLDRRNE
ļ						HVESKGKPFTMGLSPRVLSASR
						RDGLRERKGSFFRTVVLHDTKT
		İ				DSNSDTETNSSTPPRTLLEMQIL
						SPTWTYEISSSRTLGCGPSVFQQ
						RASYSPSVSGPRLGCPLMDGGR
						MRCPPGNPESCVSEANLIPISVP
						VAPHAHQHLVLSVFWKITIVIG
						GKTCVAQNHSNEAKMKQRPEI
						NHASPPGAFAWSQSGEPIASAM
	l					TVPA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15217	45585	A	15307	li .	785	MWDVGQDSSLLRGAGGPGLQL
						WFGRLQLHLGGLREHRDAWV
		ļ				HSRGLGSCSGMQGAATPTQNG
						RGSCLSPAPASSMEHAAPAVPP
						CCSWCDSSSRSRWPATATTETG
						PRTVPSAQESAVGLLTASLLKG
						DSGSTGFCPLLMSLAAPGTQQC
						SLGFCCTFERKRSSVSPAAPGTA
						APELAAALMLMPRKNQIAVYE
						LLFKQGVMVAKEGVHLPRHPE
						LADKNVPNLHIMKAMQ\SLKSQ
						GHREEQFAWRHFYWHLTREGI
		1				QSIQ
15218	45586	Α	15308	1	392	
15219	45587	Α	15309	1	716	
15220	45588	Α	15310	150	372	RRSNEWGVPVPVPARTGGG/HG
	į					QKTPPSPPLPPSSPPPSPSPPPPPA
		l				\APAKIPPPLPLPPLPPREAPPAP
		ŀ				GPALP
15221	45589	Α	15311	218	740	KSCPETVSRRKNVGSIKKHQGD
		İ				KFNPVCLKMTTKTEFVLCGWV
						PCRCL/REGEEETVTTLDYSHCS
		l				\LEQVP\KEIFTFEKTLEELYLDA
		ł				NQIEELPKQLFNCQSLHKLSLPD
İ		1				NDLTTLPASIANLINLRELDVSK
		Ì				NGIQEFPENIKNCKVLTIVEASV
						NPISKLPDGFSQLLNLTQ
15222	45590	A	15312	96	297	
15223	45591	Α	15313	469	5174	KMTSLAQQLQRLALPQSDASLL
						SRDEVASLLFDPKEAATIDRDT
						AFAIGCTGLEELLGIDPSFEQFE
						APLFSQLAKTLERSVQTKAVNK
		İ				QLDENISLFLIHLSPYFLLKPAQ
		İ				KCLEWLIHRFHIHLYNQDSLIA
		ļ				CVLPYHETRIFVRVIQLLKINNS
						KHRWFWLLPVKQSGVPLAKGT
						LITHCYKDLGFMDFICSLVTKS
						VKVFAEYPGSSAQLRVLLAFYA
						STIVSALVAAEDVSDNIIAKLFP
		<u> </u>				YIQKGLKSSL
15224	45592	Α	15314	267	495	GCLL**CLCIRL/SKNRAERALR
						PYSDVEALRLSSDFPFRQRPKG
						GSGWSLLSVEINQYYGPRVMIG
ļ	<u> </u>		l			RSNSEVVLVL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l .	*=Stop codon, /=possible nucleotide
ĺ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15225	45593	Α	15315	3	3027	AMEQAGTRPAATEHPRLRRPM
						PWLLLLPLLLLLLLLPGPAASQ
						LRYSVPEEQAPGALVGNVARA
		ĺ				LGLELRRLGPGCLRINHLGAPSP
						RYLELDLTSGALFVNERIDREA
						LCEQRPRCLLSLEVLAHNPVAV
						SAVEVEILDINDNSPRFPRPNYQ
						LQVSESVAPGARFHIESAQDPD
				:		VGANSVQTYELSPSEHFELDLK
						PLQENSKVLELVLRKGLDREQA
ļ						ALHHLVLTAVDGGIPARSGTAQ
						ISVRVLDTNDNS
15226	45594	Α	15316	1	775	MPCVNGKYNSTIQAGLQMAQI
İ						LQDEGPYEVGIDEETGSQKGDG
						IIRSSRKLLFTMALFLETNLLKD
		1				PLSRERRWRLDELDSGGREKG
						VAVFLRSAGIRQVQVAMEQAG
						TRP/AATEHPRLRRPMPWLLL/L
			ļ			PLLLLLLLLPGPAA/SQLRYSV
						PEEQAPGAL/VGNVARALGLEL
						RRLGPG/CLRINHLGAPSPRYLE/
	1					LDLTSGALFVNERIDREACLGS
	İ					AGANPVAVSAVEVEILDIN/DNS
		<u> </u>				PRFPRPNYQLQVTARCWWTSW
15227	45595	Α	15317	2	165	
15228	45596	Α	15318	1239	2251	TCSSRARRPRRGRTVSQVRGPA
						PWAPRRARLQQRGSLRGAARS
						PASPTSA/PSYSRASASAGPAAL
						QPQGAVTPGPSANLAAPGGTL
						VPAEGPRMPPVTAQSASSHPAP
			1			SASQPSSPAAASRGSLSVPQEVT
		İ				ATLLASASSPTLDATRDAPATP
1						RIAAPPGSATSSRGLGSQAQGSS
						LGTKESISDSFGLGSPAVTQRTT
			•			APPDSAALSQDTRPEAAGSWPG
						PQTGACSVGSGTDRRDTAAAT
						EAQHLSSESKEKTSAQKSGNCF
			:			GILKADFTISTLMDPEEMKDQF
						LRQVRLPSTDALKKQSISNYRP
						PGKGKPQQYLGDSAWNMVSA
	1.5.5.	<u> </u>	1.5210		0.50	GTIEGFYKE
15229	45597	A	15319	46	2859	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15230	45598	Α	15320	240	924	NGEQKCILCHIVYSSKKRH\SHE
		ļ		·		CRRCGPTEVSLSAYAKHISGQL
		l				HKDNVDAQEREDDGKGEEEEE
						DYFDKELIQLIKQRKEQSRQEPS
						SSSQEVNSDDSSSATFSVSVFYV
						WLKTILLLTWPKEVKRLDTPGL
		l				ELWLVPNMAELSEPEGPVDWK
						ERCVALESQLMKFRVQASKIRE
						LLAEKLHHNERLEYRDORKDS
						ASFQRKKQFTYKASGIEMASDL
						SKAPLEIRG
15231	45599	A	15321	3	755	LRLPRRSTRTRRGIWVRTGAAM
		l				GKVNVAKLRYMSRDDFRVLTA
						VEMGMKNHEIVPGSLIASIASL
			1			KHGGCNKVLRELVKHKLIAWE
						RTKTVQGYRLTNAGYDYLALK
				1		TLSSRQVVECVG\NQMGVGK\E
				1		SYIYIVANEEGQQFALKLHRLR
						RTSFRNLKNKRDYHIHRHNVS
		ŀ				WLYLSRLSGMKEFCYMKQFCD
						EQGSQLRIHMMGGQMWAVPIC
		İ				QPRDLFVWFVLRAEACAGCGA
						TSQQRGALGVTSTCYRD
15232	45600	Α	15322	1	1028	MNKTITTQEEVLPLQADVVQRE
						EELNSLKQKLAAALLAEQEPQP
				1		ERLVSELQL\PRKAAVFQDEILR
						SSRQLVLPELGVHRHVLVVCC
						GGLRCPLAGGLAAASVGRLGL
				1		VDCDVVEMSNLAYQVLHGEAL
						AAQAKAASAAASLRGLNLAPP
	ļ					PAETVTNCADGGMLRVLTGVL
						GCLQPLKVLKIAASLGPSYSGS
						LLLFDALRGHFCCIRLRSCRLDC
						AACGKWPTVTHLLDYEAFCGS
			-	1		SATDKCRSLQLLGPEDRVSVTD
						YKRLLDSGAPHLLLDIRPKMEL
						PRKAAVFHDEILRSSRQLVLPEL
						GVLRHLLVVGCGGLRCPLAQY
15000	45.601	ļ. —	15202	02	770	LAAAIVGRLGLVAYDVV
15233	45601	Α	15323	92	778	QGTRRWSLGTCSLMTRTQFEV
						CTKVCVQESLQRPGEELKTRCP
						SVGAGYVNYGTPAHERI/GAAV
						KTGREADKPEVTKQTQFSTRK
				1		DE/TSLFGVPLQ*LLARHRAPP*
				1		RRT/RRLLEVARTPMLPAQASR
				1		SELAARQDSAWASPAAPASPN/
		}				RSVLRRCLLSARPPLCGPL/LRV
1				1		LPSWRRQQIRENSAED*LHSEY
				1		RANRIRRRRGHPEN/WTKS*LE
15024	45600	 	15224	142	500	NPENLTRRSSSAPAPC
15234	45602	A	15324	142	500	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	ľ	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15235	45603	Α	15325	3	343	
15236 45604	45604	A	15326	1164	1648	PLMTSAPWPSCCCWRSLSSGRP AGRQRQLPTSITTANQPAPSRR KKSTSRKKQRNLPAVSPAMVAI V\RRSCDLYRWSVLHYASQERR VRDAAKPESD\RKRTTTKTRR/H
						WRYIKELESRQPGVADMRQQP TQLVEVPWNEQTPEQRQQTLQ RQHGRETRCSV
15237	45605	A	15327	28	417	
15238	45606	Α	15328	2	440	
15239	45607	В	15329	1	441	
15240	45608	A	15330	22	584	IRGRVDPPSAVEPPPKMQIFVKT LTG\KT\ITLEVEPLGYG*KNVK AQASRDK\EGI\PPDQQRL\IFAG KQ\LEDGRTLS\DYNIQK\ESTL/ HIFVLRLRGGAKKRKKKSYTT\ PKK\IKHKRKKG*SLAV\LKYYK \VDE\NGKIS\RLRRECPS\DECGC WGVYGQVTFDRHYCG\KCCLT LLFQQTRMTSNCMS
15241	45609	A	15331	3	954	LDARRHWEGTPGFTHTSPDAW ADAWGKMAGEKVEKPDTKEK KPEAKK\VDAGGKVKKGNLKA KKPKKGKP\HCSRNPVLLRGIG RYSRSAMYSRKAMYK\RKYSA AKSKVEKKKKEKVLATVTKPV GGDKNGGTRVVKLRKMPRYY PTEDVPRKLLSHGKK\PFSQHV RKLRASITPGTILIILTGRHRGKR VVFLKQLASGLLLVTGPLVLNR \VPLRRTHQKFVI\ATFTKIDISN\ VKIPKHLTDAYFKKKKLRKPRH QEGEIFDTEKEKYEITEQRKIDQ KAVDSQILPKIKAIPQLQGNLRS VFALTNGIYPHQLVF
15242	45610	A	15332	670	1008	WCILGVTPSLMSRTALFLSRHT FFA/RSHSAASKL/EKKKKEKVL ATVTKPVGGDKNGGTRVVKLR KMPRYYPTEDVPRKLLSHGKK PFSQHVRKLRPSITPG\TILIILTG\ RHRG
15243	45611	Α	15333	1	708	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15244	45612	A	15334	<u> </u>	1754	MRNEISDLTEELHQKEITIATVT
102 \		-		[KKAALLEKQLKMELEIKEKML
						AKQKPWACLVNAMPAIQVVFN
						LGRFRKCQDPHKAQEWVYPGY
		İ	<u> </u>			SGSDCEVFPEIMTSKNREEAMP
						EWKLWKKVVLSGATDAPSWD
						GPTFTDLNVFAGNMGLSDCTLS
						LGKIWKIPGHFLHTSGPVSFHT
						KGLLNTPTKERWVGGFGSEGR
				1		GVPEPGGCFGVLAVPRRTPPSA
						AAMTGGERLGRAWAGDRAAA
						SRSAPVPGTFGMMGWRPVAFG
						KNAKAYRGFGGIVTSLTPSCDK
						VTLLFVRILPGSGQSRDQSSPEK
						LTDTKSTSNKAYGFPECLRRPE
						STHIFFYDHQLGSEQPRGTKGT
						PLKWVSQKTLERETRQCQALVI
						WTDCDREGENIGFEIIHVCKAV
						KPNLQVLRARFSEITPHAVRTA
]				CENLTEPDQRVSDAVDVRQEL
j						DLRIGAAFTRFQTLRLQRIFPEV
						LAEQLISYGSCQFPTLGFVVERF
						KAIQAFVPEIFHRIKVTHDHKD
						GIVEFNWKR/LSTL*PHGLPSSL
]						SVVCGVRSLSCRSVGFCWRSTP
						DPVCLCITSGGCRTASIAEQQKL
						LPDPSYRSFIPEGQPPI
15245	45613	Α	15335	3	125	
15246	45614	A_	15336	764	1005	
15247	45615	Α	15337	1	3533	MELVNCVLMSTLGNTCCPMAI
				ļ		AMFRSSSSLPRPGHSLTGSPLRV
						NLIHIKFPLNDFHQNVVKIKNK
						DPPLGAAIGKESCPIADNELHSL
						ERRPLSPKLVQALGVAGLSRPG
						PPHTGCAIWLLLITDHKLLSEIL
						YDLLHQGLDILADPPDEGGIPR
						GKNLVELLQLPGLEEDKAFQKE
				İ		IGLKILVFKAYSTLSSDGPPLFS
1]					ALRSLQGVQSSKVGESREEKSQ
						TEEEDPGNPWVPQSLGASVSSS
L				<u> </u>		GKWGCLRTKP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15248	45616	A	15338		2800	SISGPGQEAVPLRPKAEPPEGSG MVWDRQTKMEYEWKPDEQGL QQILQLLKESQSPDTTIQRTVQQ KLEQLNQYPDFNNYLIFVLTKL KSEDEPTRSLSGLILKNNVKAH FQNFPNGVTDFIKSECLNNIGDS SPLIRATVGILITTIASKGELQN WPDLLPKLCSLLDSEDYNTCEG AFGALQKICEDSAEILDSDVLD RPLNIMIPKFLQFFKHSSPKIRSH AVACVNQFIISRTQALMLHIDSF TENLFALA
15249	45617	A	15339	8	450	TWSWSPQAPSWARPSSQARAV SLLVARGQLCQGSRRRSGVGSR VDAGAARTALRTLGTRPRLRRP CGLGSHETKAAASGQDGQRGS RGRGSLGGRELGANEPLYQ/WR AHTLTVLFILTCTLGYVTLLEET PQDTAYNTKSEIAVVHLS
15250	45618	A	15340	246	559	MPVAVMAES/AFSFKKLLDQCE NQELEAPGGIATPPVYGQLLAL YLLHN/GQRIWQRDFPGIYTTIN AHQWSETVQPIMEALRDATRR RAFALVSQAYTSIIADDFA
15251	45619	A	15341	129	915	RSGTQGRSAATAALNVIAGATP EGQSWGLAVRTVQRRGRPAK MPVAVMAESAV\SFKKLLDQC ENQELEAPGGIA\TPKVYGQLL ALYLFHNDMNNARYL*KRIPPA IKSANSELGG\IWSVGQRIWQR DFPWG/IYTTINGHQWSET\VQP IM\EALRDATRETAPLPWVSQA YTS\IIAR*FCQPLLGLPV\EEAV KGIL\EQGW\QA\DSTTRNGSCP GKPVCRGPWDVFPFNKFISLYS EP\APV\PPIPNEQQLARLTDYV AFLEN
15252	45620	A	15342]	529	MATLTISRAQTEDEADYYCHRI KLVKEGLDERTHKAYLSSSGK GCEFHMVKPGSPLGPDILGSWA QSALIQPPSVSGSPGQSVTISCT GTSSDVGSYDYVSWYQQHPGT VPKPMIYNVNTQPSRVPDRFSG SKSG\NTASMTISGLQAEDEAD YYCCSYAGSYPVVFGGGTKLT
15253	45621	A	15343	399	737	WISGGNSCATT/LSDAGLSTTSN SIGSAT\GARQAESASSQSGGA LGRHCGASGKCAESLRREPVGS ATNQWAPGWVVGRPASDSSHV VIGLGRSAWSPAEVTPSLIPGRG RTK

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15254	45622	Α	15344	1	4320	
15255	45623	Α	15345	1	966	MNVNLAEESKGKETVGDEIKG
		1				GGIQMVMLASQPLTSYCAARFI
		1				TGHGQKLKSELKKTLQVIMLNI
		ĺ				QKVRIPTRDGYNEKDILSTGKN
						VEKLELSNIAGYPSNIEKKEYQE
	ŀ					QSVLSCCSERKDANPKSVVCSF
		ļ				FMQEQCTKGEKRFLIPGTLSVD
ł						RHLRYFGILPTVSNAAVVKEVP
1						TVSNAAVVTEAPTGSNAAVVT
						EAPTGSNAARVMEVPTGSNAA
						VVTEVPTGSNAAVVTEAPTGSN
						AAVVKEAPTGSNAARVMEVPT
						GSNAAVVKEAPTGSNAARVME
						VPTGSNAAVVKEVPTGSNAAV
		l				VKEVPT/GSNAAVVKEAPTGSN
						AARVMEVPTGSNAAVVKEAPT
						GSNAARVMEVPTGSNAAVVKE
						VPTGSNAAVVKEVPTGVTLRW
		<u> </u>				SRKRPRE
15256	45624	A	15346	1	702	
15257	45625	В	15347	238	546	POORTEL KO INDIRECTOR OF THE
15258	45626	Α	15348	632	2361	FQSRTFLKNWNPGEQSQKII/CF
						QLGERAYQTDVLVKVPRQSSV
	1					FSENQRMNNPERWFESTGCGK
						TYNQNRAFNQHQRFHSGEKTY EHNECGKAFSWPSILSKHQRIH
1		Ì				TGKKLYTCEDCGKSFSVHSYFI
						QHCKIHTREKPYECIKCGKAFS
1						THSSYVQHLKIHTGEKHHECNQ
						CGKAFSHSSNLIHHQRIHSGEKP
		ļ				YKCKECGKAFNRQSNLIQHQRI
		ļ				HSGEKPYDCKECGKAFSTQLFL
						IQHQRIHTGEKPYECNECAKSFS
		į				LNRTLTVHQRIHTGEKPYRCNE
						CGKSFSQCSQVIQHKRIHTGEK
	İ	1	i			PYICNECGKSFGARLSLIQHQRI
1		İ				HTGEKPYGCREKPYECSECGK
		ļ				AFSQSFNLIHHQRTHNGEKSYE
İ		i				CNECDKAFSLLSSLVQHQRIHN
						GDKPYECHKCGKAFSQGSHLIQ
						HQRSHIGEKPYECNECGKTFGQ
						ISTLIKHERTHNGEKPYECSDCG
						KAFSQSAHLIHHQRIHTGENPY
						ECSECGKAFNVCSSLIQHHRIHT
						GEKPYECSDCGKAFSQHSQFIQ
						HQRIHTGEKPYMCNECEKSFSA
						CLSLIQHKRIHTGEKPYVCAKC

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15259	45627	А	15349	3	354	EAFGGERVSSTMSGLSGPPARR GPFPLALLLLFLLGPRLVLAISF HLPINSRKCLREEIQQGPASDW RVRDLRPVWGRWRPAQ/RTSRS QILLAIFSTPKRMQPRGNLPLTT EDYDMF
15260	45628	A	15350	283	393	YVYRRYGRFE*NDPSSALMST* GKRCQDCLDQRENKK
15261	45629	Α	15351	1	351	
15262	45630	Α	15352	1	867	
15263	45631	В	15353	1	1323	
15264	45632	Α	15354	1	3669	
15265	45634	A	15355	74	208	LESYLQLPSAEGKTVSVFGETC ATPVGP/AAGPHTQ/LAQNIVTS WLTGGRFIELKTVQILDRLELE KPCIDAEDECFNTEWSTEFTLL KAWDEYLKAWFALHLLEAMF QPSDSGKSFIFNMSVGYNLEGQ PLNPKNYPSQGVPRVLKSHRQD YLVGNKLSWADIHLVELFYYV EELDSSLISSFPLLKPHTHVDNT KKGSHPHMCAYTDYVNNPND RMPQREITFVS QRLTKLRETSRRLLSWQRSPSS TTSMHGAEWSPPRWLLAAAGV
15267	45635	A	15357	308	735	EFEEKFIKSAEDLDKVRNDGYL MFQQVP\RLRLMG*SWSRV NQRCTWAYSMTMEIFLISCVKT LSVRYAVQQLKTGVFCSLETQP SKMAPAKKGSKKKKGHS/AIIE
15269	45626	<u> </u>	15259	2	323	VVTQENTINIQKHIHEVGFKKC APRALKEIQKFAMKEMGTPVV LIDTRLNKAVWAKRIRNVTNRI HLCLSRKCNEDKE
15268	45636	A	15358	3		LSSLASMSFTTCSAFTNYWSPG SVQVPSYGTQPVSHAASVYAG LGGSGSRISVSH\SPMAGGLAG MGGIQNEKETMQSLRDRLASY LDRVRGLETENWKLESKIQEH
15269	45637	В	15359	11	991	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15270	45638	Α	15360	40	1030	RGDPRVRRRTQSLSPVLSLSPDS
						MSFTTRSTFSTNYRTLGSVQAP
1						SYGARPVSSAASVYAGAGGSG
ŀ						SRISVSRSTSFRGGMGSGGLAT
						GIAGGLAGMGGIQNEKETMQS
						LNDRLASYLDRVRSLETENRRL
						\ESKIREHLEKKGPQVRDWSHY
ŀ						FKIIEDLRAQIFANTVDNARIVL
						QIDNARLAADDFRVKYETELA
						MRQS\VEN\DIHGLRNVI\DDTN
						YHTDLQLETEIEALKEELLFMK
						KNHEEEVKGLQAQIASS\GLTV
						EV\DAPK\SQDLAKIMA\DIRAQ
		ĺ				Y\DELAR\KNREE\LDKYWSQQI
						EESTTVVTTQSAEVGAAETTLT
15271	45639	Α	15361	1	438	LHSDDLSLQMLRCSPVATAASY
						TSGPGAPISSSSFSRVGSSSFQGG
1						LGGGFGGASGIGGITAVMVNQS
						LLRPLNLEVDPNIQARF/VHTQE
				-		KEQIKTF/NNKFASFIDKVRFLE
						QQNRMLEAKWSLLQQQKMAQ
						SNMDNMFQSYINNL
15272	45640	Α	15362	176	1455	EIFPLVIQKSYKVSTSGPRAFSS
İ						R\SYTSGPGAPISSSSFSRVGSSS
		İ				FQGGLGGGFGGASGIGGITAVM
•						/VQVLHTQEKEQIKTF/NNKFAS
•						FIDKVRFLEQQNRMLEAKWSL
						LQQQKMAQSNMDNMFQSYIN
						NLRWQLETLGQEKLKLEAELG
İ						NMQELVEDFKKKYHDEINKHT
		l				EMENEFVLIEKDVDEAYKNKV
	i	}				ELESHLEGLTDEINFLRQLHEEE
		İ				IWELQSLISDTSVVLSIDSSHSLD
		1				MDNIITEVKAQYKEIANCSWAK
						ADSMYQIKYEDLQMLARKHGD
		l				NLRCTKTDISEMNQNVSWLQA
İ						EIKGLKGQRASLEATITDAEKR
						RELAIKDANTKLLELEAALQW
					}	AKQDMAQQLRVYQELMNVKL
	t					ALDIKTATYKKLLEGEESWQES
İ			:			RMQNMSIYSKTTSGYAGGLSS
}						AYGGLTSPP\LSYGLSS
15273	45641	В	15363	70	915	
15274	45642	С	15364	62	217	
15275	45643	В	15365	1	612	
	·					

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15276	45644	A	15366	[3	841	STPKMPKGKKAKGKKVAPAPA
13270	13011	1.	13300	ľ		VVKKQEAKKVVNPLFEKRPKN
						FGIGQDIQPKRDLTRFVKWPRY
						IRLQRQRAILYKRLKVPPAINQF
ŀ		ĺ				TQAL\DRQTATQLLKLAHKYRP
ŀ		İ				ETKQEKKQRL\LARAE\KKAAW
	•					PKGTFPTKRPPV\LRAGVNTVTT
		İ				LVENKKAQLVVIAHDVDPIELV
		1				VFLPAL\CRKMGVPYCIIKGKAS
						LGRIVHRKTCTTVAFTQVNSED
		l				KG\ALAK\LVEAIRTNYNDRYD
						EIRRHWGGNVLG\PKSV\ARIAK
						LEK\AKAKELATKLGLNVHC
15277	45645	Α	15367	2	1006	WINLKGNLLRKRSLTSNFWPCK
			4			LRSRKRHTTRNLRMNLWLSSH
						PPQQSNTKPWCCQLSCSTSLLT
		i	•			LLGTSLPLLSYLIHLIPPGPRQYP
						VIHRTSTLLLQHISVGKEKFHKS
						QHWGFCNNVMMLQLQLSPLK
						GLSLVDKVNRPPALSGTRVLAS
		l				KTAWRIFQEP/SEPKTKAAAPG
-						VEDEPLLRGNPCRFVIFPIEYHD
1		İ				IWQMYEKAEASFWTT\EEVDLS
		1				KDIRHWESLKPEERYFISHVLVF
		1				FAASDGIINENLVERFSQIVQITE
		1				ARCFYGFQIAMENIHSEMYSLLI
		1				DTYIKDPKEREFLFNAI/ETMPC
						VKKKADWALRWIGDKEATYG
15278	45646	Α	15368	3	1368	QAQPMGRVGGMAQPMGRAGA
İ		1				PKPMGRAGSARRGRFKGCWSE
		1				GSPVHPVPAVLSWLLALLRCAS
		1				TMLSLRVPLAPITDPQQLQLSPL
		1				KGLSLVDKENTPPALSGT\RVL
		1		i		ASKTARRIFQEPTEPKTKAAAP
						GVEDEPLLRENPRRFVIFPIEYH
ļ		1				DIWQMYKKAEASFWTAEEV\D
		1				LSKDIQHWESLKPEERYFISHVL
		1				AFFAASDGIVN\ENLVERFSQEV
		1	ļ			QITEA\RCFYGFQIAMENIHSEM
		1				YSLLIDTYIKDPKEREFLFNAI\E
						TMPCVKKKADWALRW\IGDKE
						ATYGERVVAFAAVEGIFFSGSF
						ASIFWLKKRGLMPGLTFSNELIS
						RDEG\LHCDFACLMFKHLVHKP
						SEERVREIIINAVRIEQEFLTEAL
						PVKLIGMNCTLMKQYIEFVADR
						LMLELGFSKVFRVENPFDFMEN
						ISLEGKTNFFEKRVGEYQRMGV
<u> </u>	<u> </u>	<u> </u>		L	<u> </u>	MSSPTENSFTLDADF

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15279	45647	Α	15369	2	621	RFIKMLSRHHQHVHHVLPLLRA
	1					PLPVAMEEEIPALFIDNGSGM/C
	İ					KSSFA/GDNALRAIFPSIIGHPRH
						QGVMVGMGQKDSYVGDQAQS
	1					KCGILTLKYPIKHGIVTNWDDM
						EKIWHHVFYNELCVALEEQVV
						LLTEAPLNPRANREKMTQIMFK
						TFNTQAMYVAIQAVLTLHSSGC
						TTGIVMDSGDGVTHTVPIYERH
i						TLPHTILHLDL
15280	45648	В	15370	1	966	
15281	45649	Α	15371	2	228	
15282	45650	В	15372	1	816	
15283	45651	Α	15373	256	384	RKLLNEAPC*PMR*HSLLNRLA
i						TSCNSWTVPVVPKFSKLRRGL
15284	45652	Α	15374	145	375	KIAGKNRLNRNQCKERKVQVL
				1		WISEKKKQREAVEDSPSLDCCH
İ						KFPPGS*ERLHG*KRGQWF*PP
	1					HTVEKQMRSCLL
15285	45653	Α	15375	319	482	
15286	45654	A	15376	4776	5910	VCSTAGDVNGVCLLYDLLHIA
						VASGSAKGECFFARKPLLAEGV
1		1				IYDCLQEFKEKKLVPATPHAQV
		1				LSYEALLSAHDTIAQKDFEPLLP
						PLPDNIPESEEAMRIVCLVKNQ
						QPLGATIKRHEMTGDILVARIIH
						GGLAERSGLLYAGDKLVEVNG
						VSVEGLDPEQVIHILAPRPCLSP
	}					MSLADLPPLPREVLVLAGKLPV
						PSSSFLLQPREDCQGPQPSSLIT
	İ					WFAPLETEPGVDRAMSRGTIMF
						KVVPVSDPPVNSQQMVYVRA
				-		MTEYWPQEDPDIPCMDAGLPF
				ļ		QKGDILQIVDQNDALWWQARK
				1		ISDPATCAGLVPSNHLLK/RIWT
		1				EPGICIFTKRLRCSGKQREFWW
		1			}	SQPYQPHTCLKSTLCEYCNCPT
15287	45655	A	15377	2	405	WSLVTRRALARVGLPG/SPPPRL
						LLLPLLLGWGLRVAAAASASSS
•						GAAAEDSSAMEELATEKEAEES
•						HRQDSVSLLTFILLLTLTILTIWL
						FKHRRVRFLHETGLAMIYGLEK
						ITIGVLDITGWPYCTIQPYNGFL
15288	45656	A	15378	1	3804	
15289	45657	Α	15379	349	622	RPDVANLAFLARKEGTKRGFLS
						KKTAEASRWHEKWFALYQNV
						LFYFEGEQSCRP\GMYLLEGCSS
		1				GGTSPRSFRSSSRDSPISIPLHPT
						AK
15290	45658	A	15380	314	483	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15291	45659	A	15381	352	822	DPRRVQAFPAKMKFNPFVTSDR
						SKNRKRHFNAP\SHIRRKIMSSP
						LSKELRQKYNVRSMPIRKDDEV
						QVVRGHYKGQQIGKVVQVYR
						KKYVIYIERVQREKANGTTVHV
						G\IHPSRVVITRLKLDK\DRKRIS
						DRKAKSRQVGKEKGKYKEETI
						EKMQE
15292	45660	В	15382	80	432	
15293	45661	Α	15383	11	888	VKCRKAEGRRESRLQTFEESQA
						VEAAMANVPWAEVCEKFQAA
						LALSRVELHKNPEKEPYKSKYS
						ARANM\EEVKALLGPAP\EDED
						ERPEAEDGPGAG\DHALGLPAE
						VVEPEGPVAQANRLRL\AVI\EF
	-					HLGVNHIDTEELSAGEEHLVKC
						LRLLRRYRLSHDCISLCIQAQT
						WRCIGVRDPPTSSCAFLSTSPTP
						VHSVKKIQTPGPLRPSNSTLVD
l						DMESSFQTCFASPLSQDYVNGS
						NQEEIGTGVDRCIHKFLDIARQ
						NISFLQKRLQLSVQKPEQVINED
						VDVSELRNE
15294	45662	Α	15384	1864	2705	WPGKVLAHRGLHPWHRDPHGS
						GGGAGQELRRQGGCLEQLLYD
						AAHAQRLPPLDSVLPRAALPQD
						CQRASACEGDPTLLRPSHSPGH
						PRGAEER/VPSTACLQRSWEGR/
						WNRALQQVGGLKSPWRGEYK
				}		EPRHPPPNQANYHQTL/ACPAE
						R/CFRQGPQGPGQLRRQQAEPL
						SSSLLSHQSPQSQTSLLP*L*AR
						RSLGCGNPYLCPPWSQPLPETP
						AHQSGKQPSRSRNCSSWK*NYS
		ŀ				STACPSHFLWRSRSKFSRASAST
						ASPCRMTVRRTHQRPLKARGTP
15295	45663	A	15385	1113	1378	RPHAESFVSGGIPDPPGPGPGST
						GPGPL/ERQPCSLTSWPREPPSSS
						SMKMLTQGPSLTAPYKGTILGE
						VNPAHAISSSARNTSSRTSV
15296	45664	A	15386	285	722	TPAHPGVPPGQSGRHRHW/QSA
						ARSQLQPSAW*PKTGSLFATTW
						RCQTRASTCSAHWPLMAASPG
						AGGSSMASWR/DRP*PCPPPPAP
						HLPEAAFLLR/CHDACP*NTGSA
						VPRGSCQPPAQQWDQGLAATR
						WGQAERLPRIYNWALAPP
						`
L		ــــــــــــــــــــــــــــــــــــــ		<u>. 1</u>	1	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
15297	45665	A	15387	207	605	FQASAERFGTEDD/GSLRTRPPV MQSHQWPWRRPRPLVSRGRPQ TTPERHDSGGSLPLTPRMESHS EDEDLA/EGCRWP/EAGTVGVP GPRAQGAAQR/QAVLARKKHR RRPSKRKRHWRPYLELSWAEK QQRDERQS
15298	45666	Α	15388	193	407	ACPKTSKNLRQRRLWNLAKHE F/SVDMTCGGCAEAVSRVLISLE VRVKYDIDLPNK/KVCIESEHS MDTLLATL
15299	45667	В	15389	321	625	
15300	45668	A	15390	275	690	SLWPPVIRG/SPKGNRPAILTYH DVGLNMANKKSGDHPEREGTP RSFREVVSAFGPGSSKLQIPKGS APRCGVPGRGARRELQSFRAGP RTQGSPVGPRCQSRRLRSAQEE SAKARHGGLRAHRSFSWDIRM SSKERKFS
15301	45669	Α	15391	2	137	PGRYRPN*LLLSDASPGAGNEF QSTFLTPTILPSLRFPVLYTPNS
15302	45671	A	15392		3629	MSLRKLTIMVEGKGGPNIPHAC GCRQRSSIKVSLLPAVTSKSESP KEPEQLRKLFIGGLSFETTDESL RSHFEQRRTLTDCAVMRDPNT KCSKGFGFVTYATMEEKYHTV NGHSCEARKALSKQEVASASSS QR/GRSGSGNFGGGHGGGFGG NDNFGHGENFRGHSSFGGSHG GSGYGGSGDDYSGFGNDGSNF GGGGSYNDFGNYSNQSSNFGP MTGICYTEDERSYKKNAOPTA
15303	45671	A	13393		3029	ASKKQKETQKFCLRVDGQQKV KLSVLQEKSAQLTVQLKSQKFL GHPTAGRGRSELCLDLPDPPED PVALETRSVGTWVRERDLGMP DGEAALAAKVAVLETQLKKAL QELQAAQARQADPQPQAWPPP DSPVRVDTVRVVEGPREVEVV ASTAAGAPAQRAQSLEPYGTG LRALAMPGRPESPPVFRSQEVV ETMCPVPAAATSNVHMVKKISI TERSCDGAAEMKWEDQNIGD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
]	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		•
15304	45672	Α	15394	1	1515	MGQVFRVVTGVGEQPGHPDQF
						PYIDSWLSIIQNHPKWLQACFET
						YCKTLMAQIKPGTIERDCKASV
						KEKDSQEKQKKPVLQAPPEELE
						SPALYAPIYPSLARLRQEAAPA
						APRGSDSEESTPQATPCREEPEP
						LPEKETSSSYRSNAVHFLIHNPT
						WADCKQLLQSLFNTEEHHRVI
		ŀ				QAAFQWLENNAPAGTGYIRQY
						AQQALPIEADTGWDPNQAQGL
						QSLQWYREALLNGIKAGGKKA
						TNIGKVSEVCQKPDESPSEFYER
						LCEAYQLYTPFDPEAAGNQCM
						VNVAFVSQAQGDIKRKLQKLE
						AKAVLRGKFIALNTYIKKAERA
						QTDILRSHLKELKKQEQTKSKA
						SRRKDVTKIRAELNEIETTTKI
						QKINETKSWFFEKINTINRPLAR
İ						LTKKSREKLQITSLRKEMGDTT
						TDTTEIQKIIQGYYEHLYAHKLE
						NLEEMNKFLEKYNAPSLNQEEL
-						HTLNRPIRNSEIEVVIKKLPTKK/
						SPGQDGFRAESYQIFKEALV
15305	45673	В	15395	1	1077	
15306	45674	Α	15396	69	290	YYFDSPIHPLHLSGTLRPGGSRQ
	,					FPCDPQFLCRLLHTLASL*SLPG
						DLLRGLLHKLVCRVERLDLASL
						YAVGR
15307	45675	Α	15397	62	201	
15308	45676	Α	15398	3	429	LPSSVPRQPKMQARCLSRSIRA
						VEIYESDLEEGQGLLHEFSQFVL
						DRPPPTQSRYAQELVHTCYLHT
						LASL*SLPGDLQRGLLHKLVCR
						AERLDLASHHAAVGRSELFFW
						TPYTHSPPGAGVGHPFPWHTRL
		<u> </u>				PWGVPAEGHQ
15309	45677	Α	15399	303	612	PSSSIPVYHCSLRGICIPVLGLKR
						AANVGVSLEASPPLLSTPGGLGI
						CLSCGGTRTVGSRVGSIFPW*G
1						EGTTGITGSISCKGSSDVGSNRT
			15.00-		0.00	SGVDFPRRVE
15310	45678	Α	15400	1	273	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	,
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15311	45679	A	15401	1	406	MEEGWIRLPDGRVAAPQLLGA
		-				AVVLAVQETTHRGQESLEKLL
i			1			GRYFYISPLSALAKTVRQRDFA
						DFGTTIKQDFRLLGQTSVDRLL
						QLSQGQAVKGNQLLPVSLTSQ
						CQVYKCVWNWWVLGLTDFKN
						EATDPPGVKLQTFLVSVTTHKG
						SVDPNSEKQQDLLQRAKEQSFH
						SVEGDPSRDFADFGTTIKQDFR
					Ì	LLGQTSVDRLLQLSQGQAVKG
					:	NQLLPVSLVKRKTTLAPNTQTA
				:		SPRALADSLMQLARQVSR*RAI
1					•	SCCPSH*LPSAKCISVSGIGGFL
						VSLTSRMKPRTLLE
15312	45680	В	15402	1	930	
15313	45681	A	15403	391	617	LQLQPTG/CNFLYFYVGYMKIL
						GFCFHHIVEERGGCLWYPCFYC
						NLSLLGG*RSCSHTRIVPSLLMP
				- -		KLGPGGQVC
15314	45682	A	15404	1	1860	
15315	45683	Α	15405	1	531	
15316	45684	Α	15406	2302	2691	
15317	45685	A	15407	520	2883	
15318	45686	Α	15408	3690	4727	ENCHWGCEHPCDIGSSIILSPLA
]		YWEQYHRVTYISRDTEGSIIVFP
						ALVTMLRNLFIILAGSSDPHFHT
						PMYFFLSNLSWADIGFTSATVP
						KMIVDMQSHSRVISYAGCLTQ
						MSFFVLFACIEDMLLTLMAYDR
						FVAICRPLHYPVIVNPHLCVFFV
						LVSFFLSLLDSQLHSWIVLQFTF
						FKNVEISNFVCDPSQLLNLACS
						DSVINSIFIYLDSIMFGFLPISGIL
						LSYANNVPSILRISSSDRKSKAF
						STCGSHLAVVCLFYGTGIGVYL
						TSA\VSPPPRNGVMASVMYAV
						VTPMLNLFIYSLRNRDIQSALRR
						LLSRTVESHDLLSQDLLHPFSC
1.50:0	15605	_	1,5400		2100	VGEKGQPH
15319	45687	В	15409	1	3195	
15320	45688	В	15410	1	1819	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15321	45689	Α	15411	399	1089	LLTPRTDLQEMPLDNAEIEWYT
						DGSYLRGEVFQWYDHLALIFVP
						QTGIEDVIWHIEALPNYIQKVCE
						RRNPYNQILSLQCAQHEIFLVS
						WKLVCKQLEGDDDMLVGTTA
						SNLLLESEQSTSNLNEKINHLEY
-		ŀ				EDQYKDNNFGEGNDGGILEDK
						LISN/NMGGPDSPDSVNPVE/P/M
						PTMTDQTTLVPNEEEAFALEPI
						DITVKETNAKRKRKLIVDSVKN
						LDSKTIRAQLNDY
15322	45690	Α	15412	3	809	FKDDDMLVSTTTSNLLLESEQS
		1				TSNLNEKINHLEYEDQYKVDNF
		ļ		•		GEGNDGGILDDKLISNNDGGIF
		i				DDPPALSEAGVMLPEQPAHDD
ŀ		1		}		MDEDDNVSMGGPDSPDSVDPV
			į			EPMPTMTDQTTLVPNEEEAFAL
						EPIDITVKETKAKRKRKLIVDSV
İ						KELDSKTIRAHLSDYSDIGTTLD
		ŀ				LAPPTKKLMMWKETGGVQKLF
						SLPAQPLWNNRLLKLFTRCLTP
						LVPQDLIKRRKGGEADNLDEFL
		}				KEFEHPEVPREDQQQ*HQQRD
		<u> </u>				VIDEPII
15323	45691	Α	15413	1	595	MVNRSHFRGVGEPCHQQNSHG
		İ				HLNSITETSNSPTLRAAPLQPTR
						DPSDMAVFEIDELPEGAVKPPA
						NKYPIFFFGTHETAFLGPKDLFP
						YKEYKDKFGK\SNKRKGFNEGL
						WEIENN\PGVKFTGYQAIQQQS
						SSETEGEGGNTADASSEEEGDR
						VEEDGKGKRKNEKAGSKRKKS
15224	45500	ļ.—	15414	41	705	YTSKVTKKLLIVINVICISIKHF
15324	45692	A	15414	41	785	GKGWAGGASGPANHGPAAGG
		ŀ				RSPGMLRGDQGLPGRGAAGM
		ŀ				ARPRSRE\YKAGDLVF\AKMKG
				·		YPHWPARIDELPEASVKPPTNK\
		}				YPIFFFWHPMKPAFLGPQ\DLFP
]		Y*EYRGKFGK\SNNRKGFNEGL
	1					W\EIENNPGS/IRFTGYQA\IQQQ
						SSFRTEGEGGNTADASSEEEGD
						RVERDGKG\KRKNEKAGSKRK
		1				KSYTSKKSSKQS\RKSPGDED\D
						KDCKEEENKSSSEGGDAG\NDT
L	1	<u> </u>				RNTTSDLQKTSEGT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15325	45693	Α	15415	1	1272	MWSGAGPGLAVWPQLVTVLSL
						SLSRLNCKWGLSFLACLLLGD
				-		MADAAMQHYGVNGYSLHAM
						NSLSAMYNLHQQAAQQAQHA
						PDYRPSVHALTLAERLADIILEA
						RYGSQHRKQRRSRTAFTAQQL
						EALEKTFQKTHYPDVVMRERL
						AMCTNLPEARVQVWFKNRRA
						KFRKKQRSLQKEQLQKQKEAE
						GSHGEGKAEAPTPDTQLDTEQP
						PRLPGSDPPAELHLSLSEQSASE
						SAPEDQPDREEDPRAGAEDPKA
						EKSPGADSKGLGCKRGSPKADS
						PGSLTITPVAPGGGLLGPSHSYS
						SSPL/SPLPSAGAIPPAHGGHQQ
	İ	ŀ				PGALLVLRSRGSGPCCCSGGCC
						CALPGRQHGPAGLTA/SASPTTS
						PCQQPLLP/SQGVWGSPLLPAPP
						AGLAPASATLNSKTTSIENLRLR
						AKQHAASLGLDTLPN
15326	45694	Α	15416	1	1152	
15327	45695	Α	15417	8	82	
15328	45696	Α	15418	1	1350	MGSEKDSESPRSTSLHAAAPDP
	1					KCRSGGRRRRLTLHSVFSASAR
	•					GRRARAKPQAEPPPPAAQPPPA
		l				PAPAAAQGPPPEALPAEPAAEA
	İ					*
		1				EAEAAAAAAEPGFDDEEAAEG
						*
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC ASCPKLTCEREGCQTEFCYHCK
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC ASCPKLTCEREGCQTEFCYHCK QIWHPNQTCDMARQQRAQTLR
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC ASCPKLTCEREGCQTEFCYHCK QIWHPNQTCDMARQQRAQTLR VRTKHTSGLSYGQESGPADDIK
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC ASCPKLTCEREGCQTEFCYHCK QIWHPNQTCDMARQQRAQTLR VRTKHTSGLSYGQESGPADDIK PCPRCSAYIIKMNDGSCNHMTC
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC ASCPKLTCEREGCQTEFCYHCK QIWHPNQTCDMARQQRAQTLR VRTKHTSGLSYGQESGPADDIK PCPRCSAYIIKMNDGSCNHMTC AVCGCEFCWLCMKEISDLHYL
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC ASCPKLTCEREGCQTEFCYHCK QIWHPNQTCDMARQQRAQTLR VRTKHTSGLSYGQESGPADDIK PCPRCSAYIIKMNDGSCNHMTC AVCGCEFCWLCMKEISDLHYL SPSGCTFWGKKPWSRKKKILW
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC ASCPKLTCEREGCQTEFCYHCK QIWHPNQTCDMARQQRAQTLR VRTKHTSGLSYGQESGPADDIK PCPRCSAYIIKMNDGSCNHMTC AVCGCEFCWLCMKEISDLHYL SPSGCTFWGKKPWSRKKKILW QLGTLIGAPVGISLIAGIAIPAM
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC ASCPKLTCEREGCQTEFCYHCK QIWHPNQTCDMARQQRAQTLR VRTKHTSGLSYGQESGPADDIK PCPRCSAYIIKMNDGSCNHMTC AVCGCEFCWLCMKEISDLHYL SPSGCTFWGKKPWSRKKKILW QLGTLIGAPVGISLIAGIAIPAM VIGI\PVYVGRNIHSRYDGKEN\S
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC ASCPKLTCEREGCQTEFCYHCK QIWHPNQTCDMARQQRAQTLR VRTKHTSGLSYGQESGPADDIK PCPRCSAYIIKMNDGSCNHMTC AVCGCEFCWLCMKEISDLHYL SPSGCTFWGKKPWSRKKKILW QLGTLIGAPVGISLIAGIAIPAM VIGI\PVYVGRNIHSRYDGKEN\S KHKRNLAITGGVTLSVIASPVIA
						EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC ASCPKLTCEREGCQTEFCYHCK QIWHPNQTCDMARQQRAQTLR VRTKHTSGLSYGQESGPADDIK PCPRCSAYIIKMNDGSCNHMTC AVCGCEFCWLCMKEISDLHYL SPSGCTFWGKKPWSRKKKILW QLGTLIGAPVGISLIAGIAIPAM VIGI\PVYVGRNIHSRYDGKEN\S KHKRNLAITGGVTLSVIASPVIA AVSVG\IGVP\IMLAYVYGVVPI
15329	45697	Α	15419	2	540	EAEAAAAAAEPGFDDEEAAEG GGPGAEEVECPLCLVRLPPERA PRLLSCPHRSCRDCLRHYLRLEI SESRVPISCPECSERLNPHDIRLL LADPPLMHKYEEFMLRRYLAS DPDCRWCPAPDCGYAVIAYGC ASCPKLTCEREGCQTEFCYHCK QIWHPNQTCDMARQQRAQTLR VRTKHTSGLSYGQESGPADDIK PCPRCSAYIIKMNDGSCNHMTC AVCGCEFCWLCMKEISDLHYL SPSGCTFWGKKPWSRKKKILW QLGTLIGAPVGISLIAGIAIPAM VIGI\PVYVGRNIHSRYDGKEN\S KHKRNLAITGGVTLSVIASPVIA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
15330	45698	Α	15420	1	530	SRTTPGKAPKETWGQDAGSGK
						GGQGPPTRKPPRRTSSHLPSSPA
						AGDCPILATPESPPPLAPETPDE
		1				AASVAADSDVQVP\GPAASPKP
						LGRLRPPRESKVTRRLPG\ARPD
						AGDGDHLSAVAERPKVSLHFD
		ĺ				TETDG\YFSDGEMSDSDVEAED
		1				GGVQRGPREAGAKE\VVRMGV
						LAS
15331	45699	Α	15421	1	2989	MLLVMYWAPAPPPVHMSQFLT
	ł	į				TGPKCLELSLDHGRDSQEITAS
		1				GLLWLQHAACGLAQWNQKPH
		ł				PPFLTLLQAEEPFERARSCDIYIR
						IKMLQTAQQHQVGLAPTERKE
ŀ	[ALRGSQSPQKSASGQRPGSLDR
						ELLMLIKISKCLRVPSGPSSVLG
		ŀ				TESTGPVPGELVLCQQDAGATQ
		ŀ				EGVCYGRELMICRCDSFPRLME
		ŀ				PWGVKGTPEVFRTDLITAMKIP
	ŀ					DSYQLSPDDYYILADPWRQEW
		•				EKGVQVPAGAEAIPE
15332	45700	Α	15422	218	404	
15333	45701	Α	15423	1	422	
15334	45702	Α	15424	160	470	LTMLFAACVFSGDEIMASEERQ
1				1		PMEEEWSRSSTISGEDGERWTP
	`					DTAGKAAQGIAGDVVWVSPLP
	1	l	1	}		PSKVSAASKPPKS\HHPFFKWH
						NFSNLMKMFYKMYFSKN
15335	45703	Α	15425	21	375	TFENTRFWDVQIPVPTGERRNE
						WALPPPGAGSVPAGGWSRPLTP
1						GRCDSQGRGRSRRAAPLSRSPL
]					PPLRRSPTLPGEPPPAHRPLRVH
	1			1		PRHGPRPSHL\ACLNSQAPDASR
						LRASANQ
15336	45704	Α	15426	155	340	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15337	45705	A	15427		2658	MIINTIITTTTITIIFIITIMSSSPIT ITNIITIIQHHHHHHNDQRPSLGK CRGLYIRVRKVPGERATANSRC PINAQGIESCWNGVMALWLGP NSTCSRLPELTDSEDCQPADSP GDEDPKPWWDPPHEAPLTGYF HPYGAAFVPRHAGNFKDRREV PHQDTSFCSAGQFWILPSACKT GRVALTLTLLIGAYQTLTEADV KGLTAARAASPGAKWRTQFPC PAGTCSSQAGNGQLEDCLPYPP GPSAPVGSLSRSSANGEFIIGRVI KAMNNSWHPECFRCDLCQEVL ADIGFVKNAGRHLCRPCHNRE KARGLGKYICQKCHAIIDEQPLI FKNDPYHPDHFNCANCGIKLG VFLTDRKDLTADAQELKGELY CLPCHDKMGVPICGACRRPIEG RVVNAMGKQWHVEHFVCAKC EKPFLGHRHYERKGLAYCETH YNQLFGDVCFHCNRVIEGDDE GQQLKSIRKGPLAQSQPLPTPV MLHPGCKTQIREAPGSAALESV RRPRPGLVPEEESESAAVEAV RHIHLQNFSRSLLETLNGQKLG GHFCDVTVCIHEASLRAHRCGL AIGSPSFQDKLLLGPSEIRVPSV VPVQTVRQLVELLYSGSLVVA QSEAPQVLMAASVLRIDSYRR MHADYRTRSSAPASTSAPAPLP TPVPPPLAPEQLRHRLRHLLAA
15338	45706	Α	15428	708	892	
15339	45707	A	15429	2	713	
15340	45708	A	15430	212	1246	DLPLAVHGQVSDSGAQAVASR SGDESSIVCLKEESTVNMTFRCS LPLAVISVVSFTAKCQVKSEFII GRVIKAMNNSWHPECSAVTSA RKFWQILHLSRMLVDRKDLTA DAQELKGELYCLPCHDKMGVP ICGACRRPIEGRVVNAMGKQW HVEREQYMDPGILSALIDPKYL LILSHRGLHMKSMKGFYFAKL YYEAKEYDLAKKYICTYINVQE RDPKAHRFLGLLYELEENTEKA VECYRRSVELNPTQKDLVLKIA ELLCKNDVTDGRAKYWVERA AKLFPGSPAIYKLKN/DGWVRG QMVPKPDDEGVQVRGRSGRPD AGWRLNRRSSVGGSAWRLRPQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15341	45709	A	15431	129	444	WGGPGRQPDGGRVPGRCGPRR
13341	143707	`	15451	127		PATWESPAACRPRPQSRHHSFE
						*GGSRSGSSHKRPCPSSPAVLRA
						RDQVVLHTSFEATEGDSSHEDR
			:			STPLARRCLCSHRSGEA
15342	45710	Α	15432	1	2446	METSLMVQVCGGRDCNALLPG
13342	43710	1 1	15452	1		NLCSQPTQVLPLEVDGDDGPG
			}			GLPLNAAPPSSVPPVPPSEALPP
						PACPSAPAPWRSIISRLFGTSPA
						AEAAHPPPGGALDFGALRTSSC
						LSQSHGHIPGGPGLGLGAVRVP
						VPSEEDPASLFTEPVPAAEAPAT
		İ				VOSVEDFVANDRLDRSFLEDM
		l				TPARDEKKVGAKAAQQDSDTG
						RASRAPEQQIPHGGHWRGHSRS
						YERFRPPDMVSAGPGPSDGEAL
		l				GGNPIVAGFQDDVDLEDQPRGS
				ļ		PPPPAGPVPSQDITLSSEEEAEV
						ADPPKGPAPAPQQCSEPETKWS
						SIPASKPQRGTAPMTAAASPWP
						GGASVHTGLEKCSSTRPPAEIEP
						GKGEQASSSESDPEGPIAAQML
						SFVMDDPDFESEASDTQCRAVR
						RVLPAQRAENAYGSAFQPGAL
						KPECRPTLKGGPGPVPSKAGKK
						TEATSPRPTGGASPLAHGATVG
		l				AAWGLGMASWPIGSSLAPTRA
	1					RTREPRLQRRVGSELPGDPCSG
	İ					SGASCRQVCARGRATSVRLPAL
						ESLWPLSSDGSAAAREGLEGPR
						LRQAAAPGGPGPPLPPLSRPPSP
1						ARRMRGGAAVAGRGAAAPLE
]						GRVRLGVGFLAETAVRAGAGA
1						LAAADAPPGSRTPRARPLVAAG
						SGAAGGRCPVPSRRQREAPSPL
						GGAGVDLQLECLTSEKAGVGA
1	İ	1				ATGGVLEALARAIRQKKEIEEIQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15343	45711	A	15433	26	963	FMIWAGQNRKVQDFVIQLRML
133.13		•				TPVVPPPQAAPNAQQPRAAASA
						PQTRVRAPRWPLGNVVVLGTE
						DAAGPSRPASEPPASPALPGPAL
						ELGLPNHPKQT/HRPRLAPRRSH
						PKDSCTIFKLNSFLKNIKFNNSV
						GDQVNLDYKLISGTEYDILNFW
	ŀ					NFRILAKTGCSKDGHRNPRSRP
		ŀ				TQEEGSGKPKKSLVKERSGAQE
						DLPVPPEEKLELSGTSMGPCCFS
						KTSYGIPHIIYILSYKGTTATAG
						HRWPAAYPAPLNCSVPPPETTM
						PVSDLGSPLHWASRIATPPSMW
						LCCLLAAALRVLAVSNSSAEVY
						SQS
15344	45712	Α	15434	803	1305	RSKPCRSSHNVFSWWAHSRPC
						CPWSGYSSRLCCLSCCITLGPY
						QLKNRQQRNGSYKSFGNYLCV
						GNRLLFWVWIIFQGKGRS/WLP
						RSDWVTSYKVMVSNDSHTWV
						TVKNGSGDMIFEGNSEKEIPVL
						NELPVPMVARYIRINPQSWFDN
						GSICMRMEILGCPLPGWEH
15345	45713	Α	15435	2	377	ARELLKAVWRPHKVAVMHCR
						GHQRASTLVGLGNSRADSEAR
						KAPSAPFRASVTAPLLPQAPDL
						VPTYSKEENDFLQAKGGQVME
						EG*IRLPDGRVAAVVLAVHETT
		<u> </u>				HRGQELLEKLLGWYFYISH
15346	45714	Α	15436	77	679	KMKTNQMGLKSKVVEMNATIS
						EIRLNINRKDVEYQQEILQLEAV
						WKLHKVAVMHCKGHQRASTL
						VGLGNSRTDSESPKAASAPLRA
						SVTAPLLPQAPDLVPTYSKEEK
						DFLQVEGGQVMEEGCIRLLDG
						RVAVPQLLGAAIVLAVHETTHL
						RS/ESSLGKLLGQLFSTISHFVSL
						CQNLTQRCVTCRQHNARQGPA
		<u> </u>				VPPGMQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15347	45715	A	15437	2	881	HARCTGIAARAGLLDPRHTGEA LAARSRIDLARRPHWPRGCGTG GGGGGCAGLGGAWLRGNDVA VPARALVTAAPASSSMSPPIPA AARPVSSS\PAASPSPRCGLRAG PAELRCPRRLRQARAEEPPRAG PAPGRRRPTPGICPAPLSPRLRA GGTAPSLPGVTRGSRVGRRLRP SLGGAGCRVGKRPPRLLPPGRR CSTGLPPPAGTGAPAGVCWED GDPGHELAAALLRRLGGQIRAL AGHRTAGWAQHGLSPREAVRP APAARGTLLLRRSNSRRHLQNP
15348	45716	A	15438	3	502	ANFGGTLAT GRRDSCLPGVVAPRACLHPRG QVPRLGSVGKMATPGMSWQQ HYYGGSAAKFAPSPATAQLAG HSMDYSQEMHLKMSKKIAQLT KVIYALNTKNDEHESAIQALKD AHEEEIQHILAETREKNLQYKS KVTEELDLRRKIQVLESSLEDHI KMKQQALTEFEAYKHR
15349	45717	A	15439	29	230	LQIDCKNKPAILRGPTDPLKEA DCYCRTWERPQILLPKWEGTR KPILVI*QNKAHQHAPQITLLHQ
15350	45718	A	15440	73	335	FFWGGIEEPCFVILSGSVFWFLL IWVGASASPVVGDVFGRGGSPF PTSTVGAFTVLWGVSWVLKRV CGSSWDC*FVLAINLELKFTM
15351	45719	Α	15441	373	513	
15352	45720	A	15442	1585	1649	KNSNIQHSPKITLVHQQWFQTK KKFLIYLKKNSGGLVDP*MVHT TRLCADNPQYQPRARKSHS*VK GESTTLREHPMGQKNLNNSLQP
15353	45721	В	15443	1	696	
15354	45722 45723	B A	15444 15445	113	525 474	SIPYKNKEEEEGKGGEYYIKGT CHGTKGSEQQPSALDLPSDRAY PNEKEPEN*LW*LI*QNKAL*HS PKNHTSSP/RNGSKPRKKSLINL KKNSGG*LLS*LGRHQRKVKPN ARKSKNQYKK
15356	45724	A	15446	47	614	TGCPRCKQNSTCIAAVKMEGPL SVFGDRSTGETIRSQN\DVTITN DGATILKLLEVEHPAAKVLCEL ADLQDKEVGDGTTSVVIIAAEL LKNADELVKQKIHPTSVISGYR LACKEAVFYLTENLIVNTDELG RDCLINAAKTSMSSDIIGINGDF FAN/MVVDAVLAIHARHRGQ/P RYPVTLLY*SHGES

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15357	45725	I В	15447	707	1826	
15358	45726	A	15448	108	397	
15359	45727	Ā	15449	1726	2464	SKAVQLEAGGLLQLFCCKMAP
						DPNITDVFGRTALHYAVYNEDT
						SMIEKLLSYGANIEECSEDEYPP
						LFLAVSQRKVKMVEFLLKKKA
						NINAVDYLGRSALIHAVTLGEK
						DIVILLLQHNIDVFSRDVYGKL
						AEDYASEAKNRVIFELIYEYER
						KKHEELSINSNPVSSQKQPALK
						ATSGKEDSISNIATEIKDGQKSG
						TVSSQKQPALKDTSDKNDSVSN
						TATEIKDEQKSGT\VSSQKQPAL
						KDTSDKNDSVSNTATEIKDEQK
						SGTSVFSETTGLEGYK
15360	45728	A	15450	2	675	
15361	45729	Α	15451	1	467	DHSSSPAREQNWMEN\EFDELT
						EVGFRRWVITNSSELKKHVLTQ
						CKEAKNLEKRLGELLTRITSLE
						KNINDLMELKNTARELHEAYTS
						INSWINQAEERLSEIEDQLNEIK
						REDKITEKMKNSTTVRVAASM
		<u> </u>			·	QSKLLQGVAEEGPLRLTRSASF
15362	45730	A	15452	2	632	
15363	45731	В	15453	355	894	
15364	45732	В	15454	99	716	
15365	45733	A	15455	1	1140	MEANDEROLNEL EMICSECONI
15366	45734	A	15456	1	14/0	MEVNREKQLNELEVIGSEEQNL EEEGLMIGGVAVRLVPDDIVIP
						GGVNATNGTEARDALRVKVA
						MSVTLLSGIIQYLLSALGWSYY
						TVDGVSQKNPRALGVTADQLH
						AIFTTMSDEQASFRFGFVAIYLT
						EPLVRGFTTAAAVHVFTSMLK
						YLFGVKTKRYSGIFSVVYSTVA
						VLQNVKNLNVCSLGVGLMVFG
						LLLGGKEFNERFKEKLPAPIPLE
	}					FFADHNSSPAREQKWMENEFD
						EWTEVSFRRWVITNSSELKEHI
						LTQCKEAKNLEKRLEELLTRITS
						LEKNINDLMELKNTAQEFHEA
						YTSINS/RNQTEERVSEIEDQLN
						EI*CKDKIR\EKKRMKRNEQSLQ
						EIWDCVKRPNLRLTGVPESDGE
						NGTKLENTLQDIIQENFHNLAR
						QANSQIQEIQRTPQRYSSRRATP
						RHIIIRFTKVEMKEKMLRAARE
	1					KGRVTHKGNPIRLIADLSADTL
						QARRQWEPIFNILKENFQTRISH
15367	45735	A	15457	1	1185	
15368	45736	A	15458	1	2367	

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15369	45737	A	15459	1	744	
15370	45738	В	15460	337	1515	
15371	45739	A	15461	1	1197	
15372	45740	Α	15462	110	714	HOREWAY WEEKA ARVA JORWANS
15373	45741	A	15463	142	601	HQRTKVYKFTKMRKNQPKKPE NSK/TQNPSSPSKDHNSLPAREE NWMENEFDKLTEVGFRRWVIT NSSELKEHVLTQCKEAKNPDK RLEKLLTRIISLEKNINDLVELK NTA*ELLEAYTNINSQIDQVEEK I/S/SIEDQLNEIKLEDEIREKKKK
15374	45742	A	15464	153	474	QTKSNNVNIKKQDVHTKTPSEG HQHQRPK\DHNSLPAREQNWIE NEFDELTEVGFRRWVITNSSEL KEHVLTQCKEAKNLEKRLQEL LTRITSLEKNINDLMELKNTA
15375	45743	Α	15465	1	2172	
15376	45744	Α	15466		1640	MKLMETLNQCINAGHEMTKAI AIAQFNDDSPEARKITRRWRIG EAADLVGVSSQAIRDAEKAGR LPHPDMEIRGRVEQRVGYTIEQI NHMRDVFGTRLRRAEDVFPPVI GVAAHKGGVYKTSVSVHLAQ DLALKGLRVLLVEGNDPQGTA SMYHGWVPDLHIHAEDTLLPF YLGEKDDVTYAIKPTCWPGLDI IPSCLALHRIETELMGKFDEGKL PTDPHLMLRLAIETVAHDYDVI VIDSAPNLGIGTINVVCAADVLI VPTPAELFDYTSALQFFDMLRD LLKNVDLKGFEPDHLTPAGNC YVVRRNEDCIGFFSGFYNVISL GVYLVEQQKGWEDTHSKTPSK GHQHQRSKVDKSTKMRKNWS KNAENSKNQNALSPPNDRNSSP ARAQNWMENKFDKLTEVGFR RQVITNTSELKEHILTQSKEAKN LDKRLQELLTRITSLEKNINDL MELKNTARELREAFTSFCVNCE AVSLITSLREDSHWLEDSKGHR M/RRHQE*KCEYFHPPHPGLQF YQLPESDSHSDPLWPHGGSFLS
15377	45745	В	15467	50	1603	1 QUI ESDONSDI E WI HOOSI ES

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15378	45746	A	15468	528 .	1413	QTERNSTSTSKTPSEGHQHQRP
		1				KVDKSTKMGRNQCKKA*\NSK
						SQNACCPPKDHSSLPAREQTW
		1				MENEFDELTEVGFRRWVITNSS
						KLKEHVLTQCKEAKNLEKRLE
		1				ELLTRITNLEKNINDLMELKNT
		1				ARELREAYTSINSRIDQAKERM
		1				SEIVDQLNERNDRENGTKLENT
		1				LQDITQDNFPYLARQVNIQIQET
		1			,	QRMPQRYYSRRVTPRHIIVRFA
		1				KVEMKEKMLRAAREKDTSVK
		1				QKVNKDIQELNSALHQVDLIAI
	İ					YRTLHPKSTEYTFFSAPHHTYS
						KTDHIVHSILCDSG
15379	45747	Α	15469	368	679	SSQLNKAPSLFNPVSEEFCLWL
		l				LLLQKELLTRITSLEKNTNDLV
		Ì				DLKNTA*ELHEAHTSINS*IDQV
		l		·		EERISECEDHLTEIRHSEKMKTA
		<u> </u>				LLK*DMQTRREKKK
15380	45748	Α	15470	865	2055	QTERNSININKKDNYAKTPSKG
	1	İ				HQQQRPKVDKSTKMRKNQHK
		ŀ				KA*KFPKPGMPSSPPK\DHNCSP
		ļ				ARE\QNWMEN\EFNEFDELTGA
1	1	}				GFRRWVITNSSELKEHVVTQCK
						EAKNLEKRL\QELLTRITSLEKN
	1					INDLMELKQDITQENFMKHTQ
	1					VSIAQADQGRKERV\SEIEDPTL
		ŀ				MDIKCEDKIREKK/IKKRNEQSL
]]	ŀ				QEIWGYVKRPNLRLISVPESDG
						ENGNK*ENILQDIIQENF/PFPNL
	1					TRQANIQIQETQRTLQRYPLRR
		l				ATPRHIIIRFTKVEMKEKMLRA
		1				AREKGRVTHKGKPIRLTENLSA ETL*ARRE**PILNILKERNFQPR
						ISYRAKLSFISK/GKREIKSFIDK
		1				QTLRDFVTTRPALQELLKEAVN
						MERKNQYQPLQKHTKLKRPLT
15381	45749	A	15471	1	315	MERCHY I QI EQRITICEROFUT
15382	45750	A	15472	1	2775	
15383	45751	A	15473	1	1017	
15384	45752	В	15474	1	3105	

SEQ ID NO:	SEQ ID NO:	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	noa	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	sequence		0,7,540,217	sequence	or peptide sequence	deterior, a possible nucleotide insertion,
İ						
15385	45753	Α	15475	1	1115	MKWLLLLGLVALSECIMYKVP
						LIRKKSLRRTLSERGLLKDFLK
			i			KHNLNPARKYFPQWEAPTLVD
İ					ļ	EQPLENYLDMEYFGTIGIGTPA
						QDFTVVFDTGSSNLWVPSVYCS
						SLACTNHNRFNPEDSSTYQSTS
						ETVSITYGTGSMTGILGYDTVQ
	1					SGYHHGNHRLNLKGLCSSKRSS
						TTPPLYDLGQVASPPSGSVSSC
		1				DDQSGSVVIFGGIYSSYY\TGGL
						NWVPVTVEGYW\QITVDSITMN
						GEA\IACAEGCQA\IVEPGTFFVT
		İ				GP\TSPIA\NIQSDIEA\SEELDG\D
			:			SVLICSAISS\LPDI\VFTI\NGVQY
1						PVPPS\AYILQSEGSCISGFQG\M
]				NLPTESEELWILGDVFHPASTLT
						VF\DRANNQVG\LAPVA
15386	45754	Α	15476	1	765	
15387	45755	Α	15477	470	479	PRNPSSFLQVQVQHRFLQLFVP
						FHFVFFLASLYVMVTLTTWFR*
						DPCVSGMALNCSFLKIASSSGE
						YLRPTPSYLNQVVRVTMT
15388	45756	Α	15478	1	896	CLPFGGPKPNPWDKDGGCQGR
						PQPRHLPGPGTAAQRRQQCPSE
						KSLL/CRWPSLTASTCSRLFYIL
						LHVGAS/CNLLPPAVKDSSGKG
ł						VGQDTQVLSGSGAVYRVCAGT
ļ						ATFHLLQAVLLVHLHSPTSPRA
						QLHNSFWLLKLLFLLGLCAIAF
		1				CIPDEHLFPAWHYIGICGGFAFI
		1				LLQLVLITAFAHSWNKNCNEAS
		ļ.				YLAEVFGPLWIVKVYSYEFQKP
						SLCFCCPETVEADKGQRGGAA
						RPADQETPPAPPVQVQHLFLQL
						FCLPLRLLPWLTLCHGYPYQLV
						QASAGCGCKPFAPYSL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15389	45757	Α	15479	1	1016	MERRGPGAATARGRARPGGGP
t						SVGLLATGSSLNPSFHGVARIVP
						GFIRIARPRDGSFAYESVPWQQS
						ATQPAGSLSVVTTVWGVGNAT
						QSQVLGNPMGPAGSPSGSSMM
1						PGVAGGSSALTSPQCLGQQAFA
1						EGGANKGYVQQGVYSRGGYP
						GAPGFTTGYAGGPGGLGLPSH
						AARPSTDFTQAAAAAAVAAAA
1						ATATATAT/GHRGCSPGEAEPG
İ						AEP\DGAMGAGQSFNSQFLQHG
						GPRGPSVPAGMNPTGIGGVMG
						PSGLSPLAMNPTRAAGMTPLY
						AGQRLPQHGYPGPPQAQPLPRQ
						GVKRTYSEVYPGQQYLQGAYP
						VHSGLSQFPTWQETPRHP
15390	45758	С	15480	1	585	
15391	45759	В	15481	1	885	
15392	45760	Α	15482	3	440	GPWPRPVAYLSKQLYRVSKGW
		ļ				PPGLRALAEMALLAQEADKLT
}						LRQNPNI/IGPHAVVTLMTTKG
						HH*FTNARLTYVKIPT*PLKF/G
						NTLNPTTLLPVSESPVEHNCVD
						VLDSVYSSRPNLRDHP*TSVDC
						ERYVDRSSFTNRCKVTR
15393	45761	Α	15483	3	368	
15394	45762	С	15484	109	276	
15395	45763	Α	15485	1	1710	
15396	45764	Α	15486	221	784	QNSLRSMCQKEKKMAAGVLT
						HTC\GPWPRPVAYLSKQLDRIS
						KGWPPGLRALAATALLAQEAD
						KLTL\GKT*I*RPPHAVVTLMNT
						KGHHWLTNARLTKYQSLLCEN
						PHITTEVCNTLNPTTLLLVSESL
						VEHNCVEVLDSVYSSRLNLRD
						HP*TSVDW*LHVDGSSFTNPCK
		<u> </u>				VTLKKMTSAAPVTPRS
15397	45765	В	15487	509	661	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15398	45766	A	15488	1370	2243	LGKLRLESCGLVAAVRPHSGGL
13376	43700	Λ.	13400	1370	2273	AMPLPNATRPASGRSED/AELK
						LALEHGHRSESLQGRKHRGVH
						DNSDIPQAALVGGTTMIIGHVL
						PDKETSLVDAYEKCRGLADPK
						l i
						VCCDYALHVGITWWAPKVKA
						EMETLVREKGVNSFQMFMTYK
						DLYMLRDSELYQVLHACKDIG
						AIARVHAENGELVAEGAKEAL
						DLGITGPEGIEISRPEELEAEATH
						RVITIANRTHCPIYLVNVSSISA
						GDVIAAAKMQGKVVLAETTTA
			1			HATLTGLHYYHQDWSHAAAY
						VTVPPLRLGIPPH
15399	45767	Α	15489	307	600	VSRRASSGSPVPLGSLHLQHSA
	1					PA/PSAAPKPRSPPARLSRPHLR
						GESAAPSPLSPGPASLAAAAAA
	1					RVPAQPSPAPSTTPAPRTTLAPR
						VPHRLAPAP
15400	45768	Α	15490	3	222	
15401	45769	В	15491	1	1131	
15402	45770	Α	15492	136	201	
15403	45771	Α	15493	3	556	ELPRRLVCSKLRADPGRLTPDA
						CARPGMSRYLLPLSALGTVAG
						AAVLLKDYVTGGACPSKATIPG
						KTVIVTGANTGIGKQTALELAR
						RGGNIILACRDMEKCEAAAK\D
						IRGETLNHHVNARHLDLASLKS
						IREFAAKIIEEEERVDILINNAGV
						MRCPHWTTEDGFEMQFGVNHL
						GHFLLTTWC
15404	45772	Α	15494	11	559	
15405	45773	Α	15495	1	312	MAPAADREGY WGPTTSTLDW
						CEENYSVTWYIAEF\SWLMSGF
						LPTPSSLRDLTASRWVRSLPPSR
ļ		1				SPAGRQPGPAEELPKASPCPWG
						KSLSRPFASFSASSGPS
			<u> </u>	<u> </u>	l	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15406	45774	A	15496	2	1605	WQLPHPPAAPSARQILRACQLS SVTAVAQSCLYGKQLC/GLTTG NIAGAGLLFFGGGIGGTILYAK WASHFRERVEKTIPYSDKLFEV VLGSAPYNVPLPKKSTQSGPLK ISSVSEVTKEFKQPASQLQKQK GDTPASATAGDTVLVPGVQHE ESLKTDHPAIDEGKPTPALSEEA SSSSIRERPCEEIAACLAQEEKQ EQVKTESLAKSFEDALRQTANV TLQAIAAQNTVVQAVNAHSNIL KAAMDNSEIEGKKKSAKWHTV QGALKEHRKAVDEAANALLKA KEEVQAAQFESKLVSQYHELV GQAQDDFKRQLDSITPEVLPGW KGMSVSDPADKLSTDDLNSLIA RAHRHIDQLNRELAEQKATEK QHITLALEKQTLEEKVQEQELK YEFEQNLSEKLSEQELQFHHLS QEQVHNFTLDINTAYARLRGIE HAVQSHAVAEEEARKAYQLW LSVEALKYSMKTSSAEMPAVPL GSAVEAIKANCSDNEFTKALAT
						AIPPESLTHGVYSEETLRVRFYA VQKLARRVAMID

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15407	45775	A	15497	89	2490	PKCCFFCLLSGETAAAQSCLCG
	1					KVVLRPLRPCRRYSTSGSSGLT
						TGKIAGAGLLFVGGGIGGTILY
						AKWDSHFRESVEKTIPYSDKLF
						EMVLGPAAYNVPLPKKSIQSGP
						L*ISSVSEVM\KESNRPASQ\LQK
						QK\GDTPASATAPTEAAQIISAA
						GDTLSVPAPAVQPEESLKTDHP
						EIGEGKPTPALSEEASSSSIRERP
						PEEVAARLAQQEKQEQVKIESL
						AKSLEDALRQTASVTLQAIAAQ
			:			NAAVQAVNAHSNILKAAMDNS
		1	:			EIAGEKKSAQWRTVEGALKER
						RKAVDEAADALLKAK*LLMDF
						NKEELEKMKSVIENAKKKEVA
	1					GAKPHITAAEGKLHNMIVDLD
						NVVKKVQAAQSEAKVVSQYH
		İ				ELVVQARDDFKRELDSITPEVL
						PGWKGMSVSDLADKLSTDDLN
						SLIAHAHRRIDQLNRELAEQKA
	Ì					TEKQHITLAL\EKQKL\EEKRAF\
		l				DSAVAKAFRNIHRKLKYRA*TG
		1				QKR*KEVRDAME\NEMRTQLR
		1				RQAAAHTDHLRDVLRVQEQEL
		1				KSEFEQNLSEKLSEQELQFRRLS
i						QEQVDNFT\LDINTAYARLRGIE
		1				QAVQSHAVAEEEARKAHQLW
						LSVEALKYSMKTSSAETTS\IPL
		1				GSAV\EAI\KANCSDNEFTQALT
		1				AAIP\PESLTRGVVTVKRPLGAR
		1				F\YAVQKL\ARRVAMIDETRNSL
	ļ	 				YQYFLSYLQSLLLFPPQQLKPPP
15408	45776	Α	15498	1	1212	MPKKRQALVEFEDVLGACNAV
						NYAADNQIYIAGHPAFVNYSTS
						QKISRPGDSDDSRSVNSVLLFTI
						LNPIYSITTPTRLNVFKNDQDT
						WDYTNPNLSGQGDPGSNPNKR
						QRQPPLLGDHPAEYGGPHGGY
						HSHYHDEGYGPPPPHYEGRRM
						GPPVGEYGPHADSPVIMVYGL
						DQSKMNCDRVFNVFCLYGNVE
						KVKISLKKQSPGGRPMGEEWL\
						DGYAVDRAITHLNNNFMFGQK
		1				LNV/CVGAQAREGSRGTGERK
						GGEWGPAEEHSEAEVLTHTEM
						GCGSVSKQPAIMPGQSYGLEDG
						SCSYKDFSESRNNRFSTPEQAA
						KNRIQHPSNVLHFFNAPLEVTE
						ENFFEICDELGVKRPSSVKVFSG
						KSERSSSGLLEWESKSDALETL
L	<u> </u>				<u> </u>	GFLNHYQMKNPSINLVT

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	·	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15409	45777	Α	15499	1	645	
15410	45778	Α	15500	3	677	SPPNPRAPYLFPASLQNPANWS
						PHFQYLPILSQPRSVLTAFRQQA
						QPRISMPQAQTPFRPQMERLDQ
						NIRRSNQPQHSLKPKLIQSKALT
						LFNSVKAEKDEEAAEEKFEVTR
						VVHIRGLIDGVVEADIVEALQE
						FGLISYVMVMSKKRQALVEFE
						DVLEACNAVNYTADNQIHIAG
						HPAFVNYSTSQKISRPGNSDD\R
						SVNSVLLFTILNPIYSITTDVLYT
15411	45779	Α	15501	2	1712	DEQRRRSGAMVKMAAAGGGG
						GGGRYYGGGSEGGRAPKRLKT
						DNAGDQHGGGGGGGGAGAA
						GGGGGGENYDDPHKTPASPVV
						HIRGLIDGVVEADLVEALQEFG
						PISYVVVMPKKRQ\SLV*FEDVL
						GAGNAV\NYAADNQIYIAGHPA
						\FVNYSTSQKISRPGDSDDSR\SV
			!			N\SVLLFTILNPIYSITTDVLYTI\
						CNPCGPVQRIVIFRKNGVQAMV
						EFDSVQSAQRAKASLNGADIYS
	1					GCCTLKIEYAKPTRLNVFKNDQ
	1					DTWDYTNP\NLSGQGDPGSNPN
		ŀ				KRQRQPPLLGDHPAEYGGPHG
						GYHSHYHDEGYGPPPPHYEGR
				<u> </u>		RMGPPVGGHRRGPSRYGPQYG
				•		HPPPPPPPPEYGPHADSPVLMV
				1		YGLDQSKMNGDRVFNVFCLYG
	1		:	}		NVEKVKFMKSKPGAAMVEMA
			1			DGYAVDRAITHLNNNFMFGQK
						LNVCVSKQPAIMPGQSYGLED
						GSCSYKDFSESRNNRFSTPEQA
				ł		AKNRIQHPSNVLHFFNAPLEVT
	-					EENFFEICDELGVKRPSSVKVFS
						GKSERSSSGLLEWESKSDALET
						LGFLNHYQMKNPNGPYPYTLK
15412	45780	Α	15502	1	528	NHQIRNDFTISPGVKADSRTSPI
! 	1					PQQPASSFDITEAAVSFAKDSLA
						GGVAAAISKMAVAPIERVK/RV
						PKEHGVLSWCGNLASVIRYFP\
						TQALNFTFKDKYKQIFLDGVDK
						RSQFWRYFAGNLASGGATGAT
				1		SLCFVYPLDFAHTRVAADVGK
						AGAERELRGFGDCLVKIYKSDG
						IK
15413	45781	Α	15503	1	1452	

SEQ ID	SEQ ID NO:	1	SEQ ID NO:	ì	1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	or peptide sequence	deterion, v-possible nucleonde inscritony
15414	45782	A	15504	2	991	RSAGVKAGSRPSPVLQQSASSF
13717	13702	``	13301	_		NMTDAAVSFAKDFLAGGVAA
						AISKTAVAPIERVKLL\LQVQ\H
				!		ASKQIAADKQYKGIIGCVVRIP
İ						KEQGVLSFWRGNLANVIRYFPT
l			:			QALNFAFKDKYKQIFLGGVDK
						RT\QFWRYFARNLASGGAAGA
						TSLCFVYPLDFARTRLAADVGK
						AGAEREFRGLGDCLVKIYKSDG
						IKGLYQGFNVSVQGIIIYRAAYF
						GIY\DTAKG\MLPDPK\NTHIVI\S
						W\MIAQTVHCCCPG*LPYPFDT
						RS/VRRNE*MQSGRK\GT\DIMY
						TGTL\DCWRKIARDEGGKAFFK
				:		GAWSNV\LRGMGGAF\VLVLYE
						*KSKKYT
15415	45783	A	15505	1	613	
15416	45784	Α	15506	1	1695	
15417	45785	Α	15507	131	723	LLEGKLTNRKDIHTKNPSVRH/
						RSSKTKERVSAMEDEINEMKRE
						EKFREKRVKRNEQSLQEIWDYL
						KRPNLRLIGVPESDGENGTKLE
						NTLQDIIQENFPNLARQANIQIQ
						EIQRMPQRYSLRRATPRHIIVRF
						TKVEMKEKMLRAAREKGRVT
						HKGKPIRLT/ADLSAETLQARRE WGPIFNILKEKNFQPRISYPAKL
15418	45786	Α	15508	1	3156	WGFIFNIERERNTQFRISTFARE
15419	45787	A	15509	1	879	
15420	45788	A	15510	1	804	
15421	45789	В	15511	1	1230	
15422	45790	A	15512	2	827	
15423	45791	Α	15513	1	1011	
15424	45792	Α	15514	1	1722	
15425	45793	В	15515	1	780	
15426	45794	A	15516	1	2712	
15427	45795	В	15517	1	855	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15428	45796	Α	15518	3	1263	GRTIQTKGKEV\ENFEKNLEECI
						TRITNTEKCLKELMELKTKARE
						VREECRSLRSQCDQLEERVSAM
						EDEMNEMKQEGKFREKRIKRN
		1				EQSLQEIWDYVKRPNLRLIGVP
		1				ESDVENGTKLENTLQDIIQENFP
		1				NLARQANVQIQERQRTPQRYSS
						RRATPRHIIVRFTKVEMKEKMG
						LLVPNWTNHSPLFRAILFDYKG
						FCRFGTTHQTGFSPAGANQRGP
						LAATLSGPGGEGQSAVARLTGL
						IGVPESDVEKETKLETTLQDIIQ
						ENFPILARQANVQIQEIQRTPQR
		1				YSSRRATPRHIIVRFTKVEMKE
						KMLRAAREKGRVTLKGKPIRL
		1				TADLSAETLQARREWGPIFNIL
		1				KEKNFQPRISYPAKLSFISEGEIK
		1				SFTDKQMLRDFVTTRPALKELL
		1				KEALNMERNNRYQPLQNHAK
15429	45797	Α	15519	1	1137	
15430	45798	Α	15520	1	1578	
15431	45799	Α	15521	1	1656	
15432	45800	Α	15522	1	1120	
15433	45801	В	15523	1	1541	
15434	45802	В	15524	1	1632	
15435	45803	С	15525	53	352	
15436	45804	Α	15526	1	973	MGKKQNRKTGNSKMQSASPPP
						KERSSSPATEQSWMENDFDELR
		ł				EEGFRRSNYSELREDIQTKGKE
						VENFEKNLEECITRITNTEKCLK
1						ELMELKTKARELQRVSAMEDE
						MNEMKREGKFREKRIKRNEQS
						LQEIWDYVKRPNLRLIGVPESD
						GENGTKLENTLQDIIQENFPNL
						ARQANIQIQEILRIPQRYSSRRA
						TPRHIIVRFTKVEMKEKMLRAA
ļ						REKGRVTLKGKPIRLTVDLSAE
						TLQARREWGPIFNILKEKNFQP
}			,			RISYPAKLSFISEGEIKYFTDKQ
						MLRDF\VTTRPALQELLKEALN
15427	45005		15527	1	2896	MERNNRYQPLQNHAKM
15437 15438	45805 45806	A B	15527 15528	50	658	
15438	45807	A	15529	1065	1260	RSHALALDSAGSSSPESPH*RAS
13439	43007	Γ.	13349	1003	1200	IPHTALGQNAGSWAGTAHSHH
						GPFLWSDPRHPQVVPHRTCCP
15440	45808	A	15530	96	219	EFSTKPRLSGP*SSWEMLILEPL
13440	42000	^	13330		219	APDTTIDLKLLYSDRDF
L	<u> </u>			L	<u> </u>	VI DI LIDEVET L'ADVEDE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15441	45809	A	15531	5459	6324	LLANVSQLLADISQLLANFTQL
						FSHFSQLLTYLSKLFTHLPQLLT
						HFPKLLTHQPELFSNQSQLHPNI
						T\RYSPTSPSYSPTSPNYTPTSPN
						YSPTSPSYSPTSPSYS/RDLTKLL
						PFQPTIHTTVSNLYPKLTQLQPQ
						FAQLQPNLTQV\PPTSPSYSPSSP
						EYTPTSPKYSPTSPKYSPTSPKY
						SPTSPTYSPTTPKYSPTSPTYS\Q
						PLQSTPQPLPSTHLLAPLTRPLP
				ŀ		PSTRPPAPPTRPPPPKAQPTLPLP
						LVTRPPAPPTVSQARLSARMTV
						TRRTEGTWGAAAG
15442	45810	A	15532	13	941	SVVLRPLLVVAAATATTPVAA
13.12	1.5010	1				AAATNTAKKTATVMVGEMVA
						GERGVLQMGHDITRFFYSNIQT
		ŀ		+		VINNWLLIEGHTIGIGDSIADSK
						TYQDIQNTIKKAKQDVIETFEN
						QVPRILNDARDKTGSSAQKSLS
						EYNNFKSMVVSGAKGSKINISQ
		1				GGGSQGGRGGPPISMWLLALC
						LVGLAGAQRGGGGGGGAPG\
		1				GPGLGLGSLGEERFPVVNTAYG
						RVRGVRRELNNEILGPVVQFLG
						VPYATPPLGARRFQPPEAPASW
;		1				PGVRNATTLPPACPQNLHGALP
						AIMLPVWFTDNLEAAATYGGC
						FRVPKKA
15443	45811	A	15533	207	696	TNLQEKNKQPHQKVGKRYKQT
						LLKRRHLCSQQAHEKMLIITGH
						QRNANQNHNEIPSHTS*NGDH*
						KVRKQQMLERMWRNRNAFTL
						WVGV*TSSTIVEDSMAIPQGSR
						TRNTI*LSHPITGYIPKGL*IMLL
						*RHMHTYAYCGSIHNSKDLEPT
						QRSINDRLD
15444	45812	Α	15534	1	307	
15445	45813	A	15535	1	381	LPPPHDRLWCVMFPTLCPSVLI
						VQFPPMSENMRCLVFCPCNSLL
						RMMVSSFIHVPTKDMNSGIL*S
						RPADIELFSINLVNKKIFLKGKF
				-		GDPLSLQRQLRNIVFSMPCCFPS
						FLLPWVFSQLSVKPH
15446	45814	Α	15536	134	278	DPAFLAPSPV/LMKQAPQATSG
				1		LMEPPGPSTPIVQRPRILLVIDD
		<u> </u>				AHTD
15447	45815	В	15537	1	840	
15448	45816	C	15538]1	891	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15449	45817	A	15539	1	3102	MYSHVITVCRHVKNGDILLLNR
						QPTLHRPSIQAHRARILPEEKVL
						RLHYANCKAYNADFDGDEMN
		İ				AHFPQSELGRAEAYVLACTDQ
						QYLVPKDGQPLAGLIQDHMVS
						GASMTTRGCFFTREHYMELVY
						RGLTDKVGRVKLLSPSILKPFPL
						WTGKQHINRTKDKNHMIISIDA
						EKAFDKIQQPFMLKTLNKLGID
						GTYLKIIRAIYDKPTASIILNGQK
			1			LEAFPLKTGTRQGCPLSPLLFNI
						VLEVLARAIRQEK
15450	45818	Α	15540	1	779	IPMKDHDAIKLFIGQIPRNLDEK
						DLKPLFEEFGKIYELTVLKDRFT
		l				GMHKGCAFLTYCERESALKAQ
						SALHEQKTLPGMNRPIQVKPAD
		İ				SESRGEDKKLFVGMLNKQQSE
						DDVRR\LFEAFGNIEECTILRGP
						ERQTAKGCAFVKYSS\HAEAAG
						RPSTRLHGSQTMPGAFVPVWV
ŀ						GQVSADTDKERTMR\RMQQMA
						GQMGMFNPMAIPFGAYGAYA
	1					QALMQQQAALMASVAQGGYL
						NPMAAFAAAQMQQMAALNM
						NGLAA
15451	45819	A	15541	1	1413	MEYYAAIKKDEFISFVGTWMK
				ľ		LETIILSKLSQMQKTKHRIFSLIV
						LCDITDILKEFTVAEAGGFALNE
						IMKEICKSRFSAKVSLHWKMW
						HSTASLGYHRVNVTNPDPLNLC
						LGHCWLGLRDTGQKQACGGP
						GVCPGMFRRACGGNAPCKEAP
						SVTEASSPDGSVLCLGSHVPITP
ļ						APLSSAQMVCHMVEDRRTVEL
						ILGATHNLFFSTSPLTWSLRAIP
						GNCVLFLKDYVTEPVRSQSILHI
						AGPTFKQTPNLPFLKFPEATKA
						HGPNVYQNDSTCPEPTAQVSH
						WSSVSSTLIASPVPGPEDRKLFV
						GMLNKQQSEDDVRRLFEAFGN
						IEECTILRGPDGNSKARPLASSQ
						GAPSVKYSSHAEAQAAINALH
						GSQTMPLMQQQAALMASVAQ
						GGYLNPMAAFAAAQMQQMAA
						LN\PNGLAAAPMTPT\QANGQP
						AAEEI\FANGIHPYPAQSPTAAD
L		1		<u></u>	<u></u>	PLQQAYAGVQQYAGR

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	ł .	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15452	45820	A	15542	254	883	DLRKDSPWEHRQCDKREPGNS APPAPHSWGFAVFNPRAGDVR QLCKRADTRTPSPRAPFGSAGA CRGCGRAGARSLASPKARTCIS SARRRAGGPLQLGAGSGQAKR PDR*TRREGANA*KTLHL/HSSL APSPGGVLAGFQEPPGRPQVAC CAATAAGGIRAGERR/PAAAAA AAPWAQKLRRLRLQRRGPPGC R**MVSFPRLFSGSTSR
15453	45821	A	15543	1	768	MCDFGKSTSLPASPTSSSANGR DNSTYRLKEQGGFMHSVASRD SCAESARYTDAHYAKSGYGAY TPSSYGANLAASLLEKEKLGFK PVPTSSFLTRPRTYGPSSLLDYD RGRPLLRPDITGGGKRAESQTR GTERPLGSGLSGGSGFPYGVTN NCLSYLPINAYDQGVTLTQKLD SQSDLARDFSSLRTSDSYRIDPR NLGRSPMLARTRKELCTLQGL YQTASCPEYLVDYLENYGRKG SASQVPSQAP\PSRVP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
15454	45822	A	15544	85	2249	SAEAMSGDGATEQAAEYVPEK
						VKKAEKKLEENPYDLDAWSILI
						REAQNQPIDKARKTYERLVAQF
						PSSGRFWKLYIEAEIKAKNYDK
						GEKLFQRCLMKVLHIDLWKCY
						LSYVRETKG\KLPSYKGKMAQ
						AYDFALDKIGMEIMSYQIWVD
						YINFLKGVEAVGSYAENQRITA
						VRRVYQRGCVNPMINIEQLWR
						DYNKYEEGINIHLAKKMIEDRS
	1					RDYMNARRVAKEYETVMKGL
	ŀ		}			DRNAPSVPPQNTPQEAQQVDM
						WKKYIQWEKSNPLRTEDQTLIT
						KRVMFAYEQCLLVLGHHPDIW
						YEAAQYLEQSSKLLAEKGDMN
						NAKLFSDEAANIYERAISTLLK
						KNMLLYFAYADYEESRMKYEK
	1					VHSIYNRLLAIEDIDPTLVYIQY
						MKFARRAEGIKSGRMIFKKARE
	1					DTRTRHHVYVTAALMEYYCSK
	•					DKSVAFKIFELGLKKYGDIPEY
						VLAYIDYLSHLNEDNNTRVLFE
						RVLTSGSLPPEKSGEIWARFLAF
						ESNIGDLASILKVEKRRFTAFKE
		ļ				EYEGKETALLVDRYKFMDLYP
						CSASELKALGYKDVSRAKLAAI
						IPDPVVAPSIVPVLKDEVDRKPE
						YPKPDTQQMIPFQPRHLAPPGL
						HPVPGGVFPVPPAAVVLMKLLP
						PPICFQGPFVQVDELMEIFRRCK
						IPNTVEEAVRIITGGAPELAVEG
						NGPVESNAVLTKAVKRPNEDS
15455	45823	Α	15545	12	427	SAEAMSGDGATEQAAEYVPEK
						VKKAEKKLEENPYDLDAWSI\L
						IREAQV**YRITFPSLWVQSHYL
						VLPVNNIFIILTL*MLFIGFQF*N
						QP\IDKARKTYERLVAQFPSSGR
		1				FWKLYIEAEVTILFYFFLYQYCS
						IHL
15456	45824	Α	15546	150	469	SAEAMSGDGATEQAAEYVPEK
						VKKAEKE\LEENPYDLDAWSILI
						REAQNQPIDKARKTYERLVAQF
						PSSGRFWKLYIEAEVTILFYFIFS
l _				<u>L</u>		YISIAAFTVVIENKLGT
15457	45825	Α	15547	1	834	
15458	45826	Α	15548	443	732	SFMRWQKGQLPQLCLHRESHC
						SPYPQPSFLGMEP/ADIHETTFN
						SIMKCDVDIHKGLYANAVLSSG
						TTMYPGIA\NRMQKEITALAPS
		1		·	1	MMKIKIIAPA

	n, \=possible nucleotide insertion) EIAALVIDNGSGMCKAGF DAPRAVFPSIVGRPRHQGV MGQKDSYVGDEAQSKRG KYPIEHGIVTNWDDMEKI
AGDI MVGI ILTLI WHH	DAPRAVFPSIVGRPRHQGV MGQKDSYVGDEAQSKRG
AGDI MVGI ILTLI WHH	DAPRAVFPSIVGRPRHQGV MGQKDSYVGDEAQSKRG
MVGI ILTLK WHH	MGQKDSYVGDEAQSKRG
ILTLE WHH	-
WHH'	
	TFYNELRVAPEEHPVLLTE
I I I APLN	IPKANREKMTQIMFETFNT
	YVAIQAVLSLYASGRTTGI
1 1 1 1 1 1	SGDGVTHTVPIYEG\YALP
	RLDLAGRDLTDYLMKILT
	SFTTTAEREIVRDIKEKLC
YVAI	_DFEQE\MATAA\SSSSLEK
, , , , , , , , , , , , , , , , , , , ,	PDSPRSSTNWQLSGFRLPL
RHSF	QPSFLG\MESCG\IHETTF\
NSIM	K\CDVDIRKDLY\ANTVLS
GRTP GRTP	PCTLGIADRMQKEITALA
PSTM	IKIKIIAPPERKYSVWIGGF
HPW/	AWLSHLSSKMWVQQARS
MTES	SGPLHSSTANGFLGGLLTL
	PFLWTKP
	ITFAPRNHLLTNTNTWTPD
	LVFDMRTSGASFTGETIER
	TGYVEKPNKQCRHSVTRY
	GHSDHTPSLTVATVATQIT
	GHGSHCDRNLSLATATA
	TPHHLLWPRWPLGSHSVT
	HGGHSDHTPSPTTATAVT
	RHPLRPQQSLGSHPVTHYS
	ISDCIPVTRYAHGGHSDHT
	MATAATQMALCHSLWPR
	YLVNKIRRKRCTDGVSPS PVFDTGIPWFESRYPSHLL
	SPPGYWGIAKRHQILNVS
	GKFKTNLLGYRQAVRHRI
	FRGSNPRTPANLFKTLTLIS
	ALENDFFNSPPRKTVQFG
	TEVLLKYKTAIEKRLEKLA
	IRDRAQMLKVENVQQAW
1 1 1 1 1	INKLPPARREDEDVKEIR
	EELRVSYFAQQLGKNSRA
	WSFHKAAIEAARTNIIQA
	'EAAQATERRIAADIDDSE
	RDGRVQYRVAKPGEVLA
	RVLNMVDLSDVYMTFFLP
	GTLKLGGEARLILDAAPD
	ATISFVASVAQFTPKTVET
	LKLMFRVKARIPPELLQQ
HLEY	VKTGLPGVAWSFLYSINQ
TICLE	RLDSIEAKLQALEATCKSL
EEKL	DLVTNKQHSPIQVPMVA
GSPL:	RTTQMCNKVR/CVNP*AT
	PVPQPTTQQYQGLDAGA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15461	45829	Α	15551	1	956	MSSGRRDALARGLGNSGDGAP
						HLGEHRQLEDVGGVTVTVTWT
		l				GRLQAAGCWTRSVSSTAAAGR
						REPSTRDGAALSTEPPDPPERLG
						HRGSRAGLTALGHVRTRGSSQ
		1				RGGRPEPWEHWLENGTRRTEL
	İ					VDGVSYESWRRRPCRGQLESR
	İ	ł		l		LVLENHVATDEDEPALKCORL
		1				EINCODPSIKSFLYSINOTICLRL
						DSIEAKLQALEATCKSLEEKLD
						LVTNKQHSPIQVPMVAGSPLRT
						TQMCNKVR/CAMHWLCVSCIF
		ļ				TCLPGWRPAAPDQGPAAISLCS
						LPSSSQGHREPLALGLPSALPPA
		ŀ	}			HRQRLRGSATVPIPP
15462	45830	В	15552	161	925	
15463	45831	Α	15553	31	856	GGFLHHASFFVGRGRTLLMYFS
						LLGATPKPEPEPEQVIKNYTEEL
				4		KVPPDEDCIICMEKLSAASGYS
						DVTDSKAVGPLAVGCLTKCSH
						AFHLLCLLAMYCNGNKGPEHP
						NPGKPFTARGFP\ASATFQTTP\
						GPQELVPYTPQITAWDLEGKVT
						ATTFSLEQPRCVFDGLASTNDT
						VWLVVAFSNASRGFQNPETLA
						DIPASPQLLTDGHYMTLPVSPD
	1			1		QLPCDDPMAGSGGAPVLRVGH
						DHGCHQQPFCNAPLPGPGPYR
						WVVPTGALGLGLCLG
15464	45832	Α	15554	827	1371	GSSAGEASSSLRRLSPSQPISRPL
						LPAGEKVLVVWIEDQTSHNIPL
		ŀ				SQQSKT/TLFNSMKAERGKEVT
						EEKLEARRGWFMRFKERSCFPN
						VKVQGEAANADIEAVACYPEY
						LAKITDEGGYTEQQIFSVDETA
						LYWKKMPSRTFLARRKNCHSH
						PSFSSQHLVQSAAINIKAKSSTS
						KKIMTP
15465	45833	Α	15555	799	906	FLNKLAFALRCGLALNSFLRKI
						QEPSLG\SKSGSLS
15466	45834	Α	15556	143	560	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15467	45835	A	15557	1059	1940	QQMQQEMRRKSKWMAMLGK
						WETYONS/TKGHLMKRVYKGI
						PMNVRAEVWSVLLNIQEIKAK
		l	į			NPRQYRVMKEKGKRSSEHIHQI
ŀ						DLAVSQTLRNHMFFRDRYGTK
						QQELFYILLAYSKYNPVVGYCR
		l				DLSHIAALFLLYLPEEDTFWAL
						VQLLASERHSCRVGEQLPLGLH
						AARPGDGHPGQPRHLPCVASLL
		ŀ				GAFRTSLLTVPKEPGTDPPEPKS
						DAFHPHQQKASHPPRGCPLCPG
						ATPSGSDSVQLALRNLRVLLPF
						SCSGSSCPWYPRKEALHPGGLE
						EQRGLQPLMGLVTG
15468	45836	A	15558	123	448	ALAAAVYPSHSERTGEEAKKIC
13400	13030	()	15550	123	1	SVR*LLRSAQEEI/FLRGKNSRT
						GVL*KLLQRK*RYQRALGLAM
						LSEWTISPKGPVSTGITKPMLTA
						SRVTVPRMTMSALLDFIHGA
15469	45837	A	15559	1	6387	
15470	45838	A	15560	52	3218	
15471	45839	A	15561	3602	4015	GSVESPLGQKTPRTSSASPCWPS
		l				TNLYRMELELYGPHVTSQVVK
						STRCAPIRWSAAPSILCEVRVPV
		Ì				TMWTCLPLPATGRLSMW*IWP
						RQWPCVLTSATQS*LTRCGGGT
						RAVSLAPQSHL*VCPAQSSWTS
						IILWT
15472	45840	Α	15562	932	1316	RVQDIPAKRGIPQFRPKIKGIGQ
						SYKQHYPARPWNVLAQKVV/V
		1				FEDQDALLPCLLTDPVLEAGVS
			•			LVRVRGRPLMRHTNYSFSPWH
		ĺ				GFTIHRAKFIQSQDYQCSALMG
						GRKVMSISIRLKVQKAVNRA
15473	45841	Α	15563	1	263	
15474	45842	Α	15564	66	185	
15475	45843	Α	15565	1	4217	MRLAEERAALAAENADGEPGA
						DRRLRLLGTYVAMSLRPAAGA
						WERCAGSAEAEQLLQAFLGRD
						AAEGPRPLLVVRPGPRGLAIRP
						GLEVGPESGLAGAKALFFLRTG
}		1			1	PEPPGPDSFRGAVVCGDLPAAP
						LEHLAALFSEVVLPVLANEKNR
						LNWPHMICEDVRRHAHSLQCD
		1				LSVILEQVKGKTLLPLPAGSEK
			1			MEFADSKSETVLDSIDKSVIYAI
			1			ESAVIKWSYQVQVVLKRESSQP
						LLQGENPTPK\VELEF

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15476	45844	A	15566	26	494	RPGCGEWLGNAELWGQECQL HYSPPVSPWENSSGPRFSISKTG SLIPPLLNDICQNQKPQIILSHPA RCLGLLDHFPQQWERRQAQNR AETG*GQRSAQGQTFRQRPLSV SGNCRVPPRATVSRK/PRPAPAY ALSLKSPRRRPPRTGPAPPGLVP
15477	45845	Α	15567	1	1242	
15478	45846	A	15568	114	1219	VDLPPGAPVRQSGSHGSVLRDH KTEGEARVLPLHQRVGKAGGR GATPGPHVHEDPVTICCGHSFC RP/C/ICLSWEEAQSPANCRACR EPSQKKDFKTNILLKNLVTIAR KASLWQFLSSEKQICGTHRQTK KMFCDMDKSLLCLLCSNSQEH GAHKHYPIEEAAEEHREKLLKQ MRILWKKIQENQRDLYEERRT AFLWRKRTLRLKGMTSGAQGT SQKVALGLLQQPLCHTVRAQS NSARVHKQRGGDAVSSGAGIP DPKAQRFPATLRSQAFTCTAGR RWGSGETVLHPLQRAGVRGAV AELHRQLLLSSSAAGFEECEGL CTSGPHITQGPDHQELYLSHEG VWTLFTCTGDRWKVLSGVPCP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15479	45847	A	15569	11	2694	MRMAATAWAGLQGPPLPTLCP
15477	173047	``	13307	1	2071	AVRTGLYCRDQAHAERATDVV
	ł					LLAPFCQPKTRSHGTCPPTERDP
						RGEGSTEYPGRVDGIQGWGTR
				İ		ALTGWTDRRLLCQACQTLPPR
		ŀ				HWFLPGARGWLGGSPCAHGQE
		1				SLPSQPSPILLRVESVKSRMLRR
						1
						RAQEEDSTVLIDVSPPEAEKRG
						SYGSTAHASEPGGQQAAACRA
						GSPAKPRIADFVLVWEEDLKLD
						RQQDSAARDRTDMHRTWRETF
						LDNLRAAGLCVDQQDVQDGN
		ľ				TTVHYALLSASWAVLCYYAED
						LRLKLPLQ\DYPTRPPTGRPAC/
						CAWLGIPNVLLEVVPDVPPEYY
				[SCRFRVNKLPRFLGSDNQDTFF
				-		TSTKRHQILFEILAKTPYGHEKK
						NLLGIHQLLAEGVLSAAFPLHD
						GPFKTPPEGPQAPRLNQRQVLF
		ŀ				QHWARWGKWNKYQPLDHVR
	1					RYFGEKVALYFAWLGFYTGWL
				·		LPAAVVGTLVFLVGCFLVFSDI
						PTQELCGSKDSFEMCPLCLDCP
						FWLLSSACALAQVREEAGRLF
						DHGGTVFFSLFMALWAVLLLE
						YWKRKSATLAYRWDCSDYED
						TEERPRPQFAASAPMTAPNPITG
1						EDEPYFPERSRARRMLAGSVVI
						VVMVAVVVMCLVSIILYRAIM
						AIVVSRSGNTLLAAWASRIASL
						TGSVVNLVFILILSKIYVSLAHV
						LTRWEMHRTQTKFEDAFTLKV
15480	45848	A	15570	1	924	MISQFIPENYFSGKSSATSGWA
						RAQRDFRVGAGSARPALLSGRL
						APPALGSERLSTRASSCGGLPR
						ATPRGRAPPPATRHPVPSTAQR
				:		LRSAAARQRLAGSSARGRDAL
						GKASWAPESVLEVLARAVRQE
						KEIKGILLGKEEVKLSLFAGDMI
						VYLENPIISAQNLLKLMSNFSK
						VSGYKISVQKSHAFVYTNNRQS
						ESQIMSELPFTVATKRIKYLGIQ
						LTRDVKDLFKENYKPLLNEIQE
						DTNKWKNIPCSWVGRINIVKM
						AILPKVIYRFNAIPIKLPMTFFTE
						LEKTK\FIWNQK\RAHIAKTIL
15401	45849	A	15571	1	825	ECKTAUTWINQK/KAITIAKTIL
15481 15482	45850	A	15572	1	1488	
13404	الاقادار	14	13312	L <u>.</u>	11700	I

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
:				sequence		
15492	45051	 A	15573	<u> </u>	1476	 MPIKVDWIKKSRTHTPRDTMSP
15483	45851	A	13373	Ţ	1470	
				-		KRGINISIQEYRNTPDASILSELR
						LERDTNTQVYTYEQEHTNKNN
						GHQAIDKQNKGHIRRESDRGEY
						YPLDCRPTGSLTDKGWGMLLC
						GMMTAGKPTPILTFLLDVGLPG
						GTAINKSSVIRVRNREEGNVEK
						WGERQVDQRDAVMRVRCGIR
						NNVEAGLKYGPGTMKVSLSVF
						YTTQQCRRYLSVLCFAGLARAS
						VPFNEDFQMSLDEKLFSNLHPL
						GSSSSPPKSLLHCPLLEHQFTML
						GFSDLGAQLASPSGSHTRAAGV
		İ				AACQPAPTLTWLSPWVARGLG
						TLEQGVVLVGEARAAQEPMEW
						VGGSCMVGCRSRALPRGKAAK
}						AWREIQRSAGAGGAPVSPWSG
						GAPSRACPPSASSVRFLCAARR
						LACNTYSPVLYCVRNWWILGL
						TDFNNEATDPRDPFLANVPGQL
						AAVAIGTCDGNVYSAGDRDYR
						FALESILKACTLALALEEVGPPA
						VQDKIGADP\TGLPF
15484	45852	A	15574	140	448	GFLGACAGREFKMAPP*SSLSG
13 13 1	15052	` `	15571			*KLSVSPCPGLLISAEGPEAVFSI
						V*NARVRGFILEVSETKNPPIPD
						TISTKHASLMSDSYREKKLVPV
						VYPHSSTPSFQ
15485	45853	Α	15575	1	1001	MEAPQGALEQDGPAVLSQQRH
113463	43633	**	13373		1001	RKKPFTATPIDPSYHRSHAISLS
						VIIGYCVPPSHAAEMAAYGWT
						i e
						MGKSKLMAGQWITQSQVFCDS STDGPRQHKAQTAILTKLQSQQ
						, , , , , , , , , , , , , , , , , , , ,
			:			RSTNNSTVSHRQQWVNCRLLR
				;		VKCQRLRAHKELLPVQRWLWR
						QGRCVPTTSLVAVSETAGAGA
						AALALSRLCRCLQVSASARPFS
						SRTLSPQPKCWAMAASHQPIKG
						ILKNKTSRTSSMVALSEQPCRT
						VHEELSKKCQKWDEMNILSTY
}						HPADKDYGLMKIDEPSPPYHG\
ļ	!					LTGDDENACSDTETTETMVSDI
}						LAKKLAAAEGLEPKYWVQEQE
		<u> </u>				SSGEEDLLAEE
15486	45854	A	15576	1	1851	OVER DE LA COMPTE DEL LA COMPTE DE LA COMPTE
15487	45855	Α	15577	73	418	GKPDGDKVSQHLLLWKG/YSF/
						PPSLMKLSLAGYEILS*KFFSLR
						TLNIGPPLSSGFRVSAERSTVSL
						MDFPLCITQCFSPAALNIFSFIST
						LVNLTIMCLGVALLEQYLCCVL
15488	45856	Α	15578	2	460	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	1100	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	sequence		05/510,217	sequence	or peptide sequence	and the second s
15400	145055	<u> </u>	115570	<u> </u>	local	NATEVEL IDVALLOCGI AGVIDG
15489	45857	Α	15579	1	2031	MATEVFLIPYALLQCSLASYPS
						KPGIYVPSPQMWVGSVTCSDQ
						KNTIEVLVPKAYTLARQQYKSL
		1	•			SEQVYSCHQGTLNTRAQESAAS
		1				LLKPARPRTRQKKETPNTSEHQ
					}	KEQTLDTPPLRTVTLTTRVRGFI
						LEVSETKNPPISDTMGLDISKQE
						EVLGRESDEQGEQSQGLSEMQ
						TALAMLLTREPISAKQAPHEPP
						RRQEDNTQAREGETRSRKTRA
						EGQKGHAHAENERNAKERESE
						RKRQEAEDENGKEKQERNEGT
						HARHMKRRKRDEKKRQGQNK
						PGNKRPEGNEEAQRRRKKNRE
						RQRFSFYQSGPSAAGLLVFAGG
						PLQSLFAWVSPLEAAEQQRFLS
			,			VPSSGPLRKFCLYFGFVGNATP
		İ				LRGLNTHGRLLTPAHGTRRRHT
						RPLNSPVPEKAGSTTESNGSPY
						GQNPGPLPPKGNRAQGEKWM
						GRRIQGTTRPRAREANHKKMT
						QIMFETFNTPAVYMAIQAMLSL
						YTYGCTTGIVMDSSDGVTHTVP
						IYEGYALPRAILCLDVAGRDLT
						NYLMKILTEHSYSFTTMAEREI
						VCDIKEKLYYVTLDFEQEMAT
						`
						AASSSSLEKSYELSDGQVITIGN ELSCCPEALFQPSFLSMESCDIH
						` i
						KTTFNSIMKCDMDIRKDLYTNT
					1	VLFGGNNMYPGIANRVQKIVTL
15400	45050	<u> </u>	1.5500	405	000	VWMGSSILALLSTFQQMWVSK\
15490	45858	Α	15580	485	886	VPYFHFTCSHSSASQLPFLQMM
						KIAKTCSLGLNRLPEAKLKAFP
						TPRVTSL/WPRQRGRPRR\GCGR
						NRSGSLVSAARGVGGRSQAAA
						ARRDAPSPPATSRMPPRQGSVS
						WGCARNAPGRRADPEVRRQPG
						PSCT
15491	45859	A	15581	1	435	HD COTC A FOTA A CNON A DODDO
15492	45860	Α	15582	2	339	HRCSTGAFSTAASNSNADRPPP
						ARRRGLPR/RGPP/SRRRSPRPPR
						REPWPRTRRPRTRTAKARRRAT
						LEGRCRRACGHWQPRAAAAA
						AAAAAVAAAAAAAAAAAA
		L				GASATW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15493	45861	A	15583	1067	2093	RRNTSSKCRSRNCRCPRSPHRH
	Ì					RPSLRPRSQQPKCKCRPRSRRSS
						RAPSSRRSRPQGLVPC*RAPPW
						PTSRWPGSPGFPLLNCSRKGRC
						RPRHPSQPRCPCRSLRWCPSRQ
			İ			LWSPHRESPPCP*TSRGSAWRS
						VSHRRQQDRPWWTQPVHMQQ
						LLKLKQQAVQQKAIQPQAAQ
						GPAAVQQKITAQQITTPGAQQK
						VAYAAQRALKNPVSYHTHLPG
:						PETGRGPASADPDPGCKTSSSCS
						TANTRGQHPASCLCFPAGFSTD
			1			CGAHAGDGGRAAGADDPCSDR
						DCPGGSAETHSAAGGDHGVGP
						APDSRRSQPSPGARQLRQPKPA
15404	150.60		15504	1	107	AQVTDEGPCCQAKDTY
15494	45862	Α	15584	10	607	RALCGESRLPLARKLS*AKKSV
						QRSTLQSRRIWVCWFHFAN*SL
						F*KQGRT*ESPL*L*LIPAS*RPS
						TSESPPL*KANFQQPHRGL*EKV
						AEGRRGP**EY\YTSSSSSSSSSS
ļ						SSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSS
		ŀ				SNNSSNSRHRRRRRRRLLSFLSR
						RRRHSRCCLSRSRPRRRPRRHP
			ļ			ARLWLRSRCTDQRKNFQLPRK
						TV
15495	45863	A	15585	217	1012	AKNDSKRQNFGHRGQPTFGAD
						KAKGERVRTSSTIRRTSYLDTIT
						GPYLTGQWPRDPHGHYPSCMK
						DKATQTP\SCWAEEGAEKRSHQ
						RSA\SWGSADQLKEETPTDLQQ
						NAQQKLYKPEEVGCLISDFLKE
						KENCQPRILYAGKLSSECTPDE
						EAVNIVEMITKDLEYYINLVDK
						EWQGLRGLTPILKEVLRVKCY
						QTASHATKKSGMKGKVSQFGK
						LHCYFKKLPQLPLPSATAILISQ
						LTSTRQELHHKKDCNLLKTQMI
15496	45864	Α	15586	2	387	
15497	45865	A	15587	1	165	LFMFFCLLYIR*QLLHPGYNAPF
						QNTAATRGRGIAQTAPEAAFQL
}						FLHQRVRFHC
15498	45866	A	15588	172	359	PDGDGIESINYLGQYGHFHDIDS
13 176	1.5300	J.,		1		SYP*AWNVLPFVCILFYFIEQWF
						IVLLEEVFHIPCKLDS
15499	45867	В	15589	1	2850	T. DDDD VI IIII OKDDO
15500	45868	C	15590	1	2835	
15501	45869	C	15591	78	491	
13301	173007	1_	1,2371	1,0		<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<u> </u>				
15502	45870	Α	15592]1	5721	MAVLSLPLAPPTLMPPAHFNLV
						EMQVKATLKEYWQRKSPGIPA
						GANRKKKINGSSPDTATSGGYH
]	1				SPGDSATGIYGEGHASSATLED
						LELRQPTPSTSPNDCSLYPSPQS
						SSNSSSLHAPHSQYRELAVALH
		i				SSSAIISQLSENINSLNNENKSAL
	}					QLERQVKELKLGELKETVTSDP
						SNNGWEAVTSLWEGERLPARG
				·		AEEPRGSESTAAARPVPVSPAA
						VRGRLVAGEPGSYQPAETAAK
						1
15502	45071	C	15502	170	1207	AQLSLMALPGEGY
15503	45871	C	15593	178	1287	
15504	45872	В	15594	52	855	
15505	45873	C_	15595	1	1698	NI ADDIDOCOCI EL IOVI I HACIO
15506	45874	Α	15596	302	617	NLAVNPSGPGLFLIGKLLIIASIS
						EHVIGLFRDSTSSWFSLGRVYV
						SRNLSVSSRFSLVFDDGDVQMG
]	1					FWCGCPLCLLVFLLT\AGPSAA
						GLLEFAGGPLQTLFA
15507	45875	Α	15597	1	2762	MYGTGYRDVAGKWAIDPDSK
1						REFLDVTQEGIQGCDFSGTCRQ
						TLSILTQPLRQWGLEGIKKPNS
				ļ		WIISEESVSNGGPPLLIPRQTAS
	1					GVDLQQTPTDLQLRVLTVRRK
						TNKQKGIASTSTKRTSTPKPHL
						YVTIIKDQSYIKPQRWGKNIAE
	-					KLKILKIRVALSLQRNAAPHQQ
						WNKAGRRMSLMSSQKKASEVI
	1					ESQMNEIKGEEKFREKRVKRNE
						QSLQEIWDYVKRPDLRLIGVPD
						SDGENGTRLENTLQDI
15508	45876	В	15598	1	1989	
15509	45877	В	15599	1	762	
15510	45878	A	15600	686	850	GQPERKVGLPTKGSPSD*QRISR
13310	1.5075	-	15000			QKLYKPKESGGQYSTFLKKRIF
						NPEFHIQPN
15511	45879	A	15601	2	382	11111/111
15512	45880	A	15602	1	993	
15512	45881		15603	3	460	SPCWGQLLQATQP/ATAFAQKD
212213	42001	A	1.5003]	1400	LLVGAVLFGHNGSLSSRNTLAF
						l .
		1		1		SMDVGPSQWMWGLLNGCGAS
		1				AHRQCTRCTLQVTIALEGPLAP
				1		LKLAWHKKIYIFLGEAAEIRRD
		1	1	1		QLEAERELFHSRAEQGRLFVGR
						PGQAAHLMLPPGDCVSIWTQH
						MIT
15514	45882	Α	15604	1	130	
15515	45883	В	15605	1	561	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	'			sequence		
	<u> </u>	<u> </u>				
15516	45884	Α	15606	94	420	EKAVGRWIFQSKRYRSPCEKPY
	Ì			1		SKGLQVQRTVPEGPSKAARAA
						QAPGPNLLPCLPT/PQTPEFIISEP
						LANMYSCGNQNTLMEELAEQA
						QQHDEMLHMHHALKEALSIIS
15517	45885	В	15607	46	747	
15518	45886	Α	15608	186	364	AFAPAPCTLVPGVLRGPPILLRE
					:	TGLAH*CPAVISPPKSLPWLPTA
						YSVESQQVILFGL
15519	45887	Α	15609	155	511	GAAVAGTQNLCLGGGGAVGG
						S*RGALNGEGLLHPGYNAPFQN
						TAATRORGIAQTAPEAAFQLFL
						HQRVRFHC*FLLWREGRGR*AR
ŀ						REPLSRPQPPAGPLGGALPAPPL
						PDASTLVGVG
15520	45888	A	15610	1352	1684	MSIRSAWCRAEFNSWISLLTFC
13320	43000	A	13010	1332	1004	LVDLSLVFDDGDIQMGFWCGC
						_
						PFCLLVFLLT\AGPSAAGLLEFA
		1				GGPLQTLFAWASAVVAAEQQI
		<u> </u>				LVSSKCCCLIVPLEVLSQRSTWP
15521	45889	Α	15611	98	367	
15522	45890	Α	15612	426	738	VCVWGDPPGPAGGHGEAAATL
						WKEKVDLKERVEKLELQFIHLS
	1					GQTDTIGRKYISQGAVSETQHW
						ERRTSSGWP\GPGGDEGEPAGA
						AGQVLQLVGDHKEGHGKF
15523	45891	В	15613	86	1411	
15524	45892	A	15614	950	1493	GTVFLWRRRGTLIFRVSGFSAL
	ĺ	1				FFPPSLWFYLPFVFDDGDLQME
	1					FWYELSFLLMLLLFLCACWFSF
	-					*QSGPSAAGLLEFAGGPLQTLF
						AWVSAAVAEEQWILVNRKCCC
						LIVPLEVLSQRSTRPCEVSVHPY
	1				1	VHCCLGIEELNIYCSLHSLGLFE
						AILLGKAFEELGCCDPSLFATV
15525	45893	Α	15615	1	5731	MDSHRPTAWPAAVLAGRSPVA
						AESERYCSRQPPLPRRQALLSV
						LVLASDTRLPFPWSSQTFCGILS
						KPLVLSGPLQTQGETNNENGLV
						GGGEVGMQPLVLFERSLFPGRC
						CVLERRIVPWMRQESYSSSPI
						WFVDSDELNLTSVLECLEDDK
						NNNLLHQQLKWLICGLCRLYN
			1			LPKHLDVAMPDQPLPMGQSGP
1						SAAGLLEFTGGPLQTLFAWVSA
1				i.		-
						ALAAEKQILVNCKCCCLIVPLE
15-5-	45004	 	15616		501	VLSQRSTWQYELKEYW
15526	45894	Α	15616	2	521	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15527	45895	A	15617	1	1261	MSEHNITEECQQENQSIEQWRSI
						AHSDSIEVSWLHEVIKTYLKAK
		1				STKLDLMKKLKELSSQPSRKPK
						PFIPSQEGTGDNDVIYSTERLSK
						LLNQKKGSTRLYDKPLSNLLNQ
	·					KKVSTRTEGLTVDSRADMRNV
						TTILLSTESSSTDSSSSMDSSSSM
						DSSSSIGRVETATCAMIPVQKSK
						RSGPPQGPSAATGDGAEGRFLP
						RCSRPWADRLAWSQPRRHLAP
						GPSAATGDGAEGRFPPLCSCCG
		1				TDRLAWPQPQGHLALLPRCGE
						CGRARELPGDCCLHAEGGCSSV
		ļ				LEVLARAIRQKKEIEEIQIGKQE
						VKLSLVADDIIFCLENATDSSRN
						LLELIKEFSKVSRYKINGHKSVA
				İ		LPYINSYQAENHIKNSTPFTIAA
						TNNNNKK\KNLGIYLAKESKDS
						YNENYKTATERNHRWSQAR
15528	45896	В	15618	60	499	
15529	45897	Α	15619	134	1416	RTSICWTMDSSVQNTVLGPTRS
						SVKGVMEWLQYLQMERR/CTL
						SISMESMPRVAEDAGDVGAAE
		ĺ				PLHLEPPAGPLEVEQQLQAEIK
						HLRKTLESLAGQLQAQVEANE
						GSSHLNRQQDERLREWKAELW
						EEQGETGRSSVQKDRTTISHAL
						SQNHQLEEQLAKERLEATSQQ
						NQQLQSQMSLMALPGEGDELG
						SKEEEEAPQPMPSILEDLKSREA
				,		MVAFLNSALVSAEEEQQAQLH
1						GQLKEQRVCCQHLAHPVALAQ
ŧ						KEPEAAAPTPRTGGDSAVWTV
]					DLQVEKTKFIMFTIKPGEVVNK
						DMAQSIYRSGKNLDKDMYGH
	1					DLEARRKTNRPLETTFYFLSYE
						LDYSRCFIGCLKCLCLPSPSATI
	1					GSFLKPSQKQKPVCLLYSLQNH
						EPIKPLFFINYPDSDSYRHTQVIR
						ILLSGSIHVQVPATKTPL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15530	45898	A	15620	1	1665	MGGHVGGRHRMLTPKLIPETL
						AAGSMKINPEEPNRPPINGAPG
						GPALPLAAGDSGSEQAGEPALA
						ASELQGWDSTLGSPGSLRGTGI
						VATRSGSRGFRLSPASLGRRSR
						DRKTKELAFAGSGLATRGQRA
						SSDAAAGPPLSRCSPARTPQLPV
						RGSLTGPRAPAPPRARVCLLSC
						SCROPFPPLTGGLNVGRPSGPS
						WKICPRSLPSASAETGIRRRRAA
						AHPPLQQPPLCGESLPSVQCSA
1						1 7
						NADSSIPVPSRWSQDACRNTGA
						AGPPGYPACVNNRNRTDKNSQ
l						LDETHHMKTKVTDRGGFGAFT
		ļ				MLLRVVDTGHTSCSGSVEWDP
	1				,	RLYGDQADLSRSGHLDRPGAH
						HDYYTAIKKCEEGKSLDTTVK
	1					VKAGVNGFGRIEHLVTKPAFNS
						GĶVDIVTINDPFVDLNNMVYM
						FQYDSTHGKFCGTVKAENRKL
		1				VASGNPITVFQ/ERDPSKIKW\YI
						AESTCIFTTMEKAGADLQGGA
						KRVIISAPSADAPMFVMDVNHE
						KYDNSLKIVSNASCITNCLAPL
						ANVIHNFGIVEGPMTTVHAITA
						TQKIVDGPSGKLA
15531	45899	Α	15621	2	1105	GRVGCSTVSRIFSCVASRATSLR
						TPMGKVKVGVNGFGRIGRLVT
						RAAFNSGKVDIVAINDPFIDLN
						YMVYMFQYDSTHGKFHGTVK
						AENGKLVINGNPITI\FQER\YPS
						KIKWGDAGAEYFVESTGVFTT
						MEKAGAHL\QGGAKRVIISAPS
						ADAPMFVMGVNHEKYDNSLKI
						ISN\ASCTTNCLT\PLAKVIHDNF
						GIVEGLMTTVHAITATQKT\LD
						GPSGKLWRDGRGAF\QNI\IPAS
						TGAAKAVGKVIP*AERGKLTG\
						MAFRVPTANVS\VVDLTCRLEK
						PAKYDDIKK\VVKQGVGGPPQG
		İ				AILGYTEH\QV\VSSDCNSGPPV
	1					FPPFDAG\AGIALNDHFVKLISW
						YDNEFGYSNRVVDLMAHMAS
15532	45900	Α	15622	121	221	
15533	45901	A	15623	189	538	
13333	1.0701	ı '' —	10023	1-37	1	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
15534	45902	Α	15624	3	1015	DRAGPPGAPRLGQQLLCAIILL
						GLAAKKKEINITIIKQLLKKFLN
						KKPSYGLERQMETTQNLVDSY
						MAIVNKTVWDLMVGVMPKTI
			}			MHVMINNTKEFIFSELLSNLYS
		1				RGDQKTLMEESAEQAQWRDE
			1			MLRMYHVLKEALGIIGDINTTTI
		1				ST\PWGPVDNSWLQVQSVLARR
						R/SPGLAPTAPSLPTPHRLSLGA
		1				LGTGSVPTLLHHPGSSSVSSLSG
ļ		ŀ				LVA/DCFGLPKARGGPGQRLPF
		ł				PSLPGRLFCDLAVAPPAPKAAP
		1				FICDFVYCCDQASEPVGCQGQT
						APHRLHGSPLGIPQAPPGSIQA
		1				WLRAAVFWKGALCPRCGRHIQ
15535	45903	A	15625	547	1242	EFLLHRCTLVSL PKALIKTSNKSSLSYQLTTHSGT
13333	43903	^	13023	347	1242	MACPWIRHEVLLSRLGFLVLEN
		1				TNMDKRPSKLLRWLERQMETT
						QNLVDSYMAIVNKTVWDLMV
						GVMPKTIMHVMINNTKEFIFSE
		1				LLSNLYSRGDQKTLMEESAEQ
						AQWRDEMLRMYHVLKEALGII
						GNINTTTIST\PWGPVDNSCLQV
						QSVLAGCSFIILVQAVFLLYQA
		1				WWLLFWAPQGERWPWTSGLE
		1				DTVTREEGSPKGLSISL
15536	45904	Α	15626	108	308	AQKHHLT*LSFLPRH*ILT*NLT
		1				KIPTSKIKKTPRSRSPRRSTSCFL
						DNLAPSCGCSGKGSSWRIC
15537	45905	Α	15627	687	1910	KDHQLRKKLRRRGKQGAFTAA
						LPPGSPRPPLRGGRSGGVGLGV
		1				RGQRTPQAPQAVTPSEGPEPLW
		1	ļ			GPVLGGGGEHRRGTGASAAAG
						ASMRRLGAAGGGAGAGPQRG
						VGARVVVSEEANSPGAAITSNR
						ETQDANEGLSAPRPECLLHPTC
						RLLYVPLIVRDWVCHTRSVVLL
						QVVNVTGIAVGTGLASAYDTL
		1				MSQSFGVKNLKRVGIILQRGVL ILMLCCFPCWADLVNTERILLL
				İ		
						LKQDPKSPGWTRECFQEWGSYI
	1					HLVIPSMFMVCTEQWTFEIGNF LAGLIDVMELGTQGIICELASV
						AYMNYFRLNPQPPISDFTINLAF
						LTLTERKILGYIQLRKGPDIVGP
				1		YGLLQPFTDAVK\FTKEPLWPS
						TSTITLYIIAPTLALSIIVLLALPT
1						RMLAATPRPRGT
15538	45906	A	15628	1	1173	THE PART OF THE PA
15539	45907	A	15629	3	1447	
	1	<u> </u>		I	<u> </u>	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15540	45908	A	15630	1	215	MDPKQEEIPDSPEKEFRRYRIAE WIRIHQPIICCLQETHLTHKDSH KLKV/KGWNKAFHANGHQKSA ARETK
15541	45909	Α	15631	1	1524	
15542	45910	A	15632	1	575	MALSERSQPQDHVSHDRIYVK HPEYANLYRQKVDYTLPRVSIS IHVGQAGVQIGNARWELYCLE LAIQPDGQMPSDKTIGGGDDSF NMFFSEIGAGKHVPRAVFVDLE PTVVDRCLGHWMAAFLGE\EV HTGTYRQLFHPEQLISGKEDAA NNYSRGHYTIGK/EDLVLDRIH KLLTSQCRKPKATMWLSQAT
15543	45911	Α	15633	208	465	
15544	45912	A	15634	107	1218	EARRPCTLHLRSALHPATMCEL YSKRDTLGLRKKHIGPSCKVFF ASDPIKIVRAQRQYMFDENGEQ YLDCINNVAHVGHCHP\EVVKA ALKQMELLNTNSRFLHDNIVEY AKRLSATLPGETLCLLFYKFRG YWPSSTNPSSRPAAPASPPAAE RAGDFRAGRKPSVPGSRSVSGS PRGPCSRLSLSTFCTRYCPLSAL TSGPGCPYPMCFFLSPRVSRLLY SSHMVAGCRALRRCKNSH
15545	45913	A	19033	3	1216	GGLWEKMAAAA\QSRVVRYLS MSRSAITA\IATSVCHGPPCRQL HHALMPHGKGGRSSVSGIVAT VFGATGFLG\RYVVNHLGTAW GSQ\VIIP\YRV*IKY*QSMHLSFP WG*PGGQLSVFLGNWGTARD* RFLFRRVSYNTGNVGSSILFGTR LGETKNF\DFWRDVFVKIPQAIA \QLSKGSWEFEKFISCFHIWNGG II*KALS*DI*GNKAVGEKVVRD AFP\EAIIVKPSDIFGREDRF\LNS F\ASMHRFG\PIPLG\SLGWKTV KQPVYVVDVSKGIVNAVKDPD ANGKSFAFVGPSRYL\LFHLVK YIFAVAHRLFLPFPLPLFAYRW VARVFEISPFEPWITRDKVERM HITDMKLPHLPGLEDLGIQATP LELKAIEVLRRHRTYRWLSAEI EDVKPAKTVNI

Sequence 09/540,217 codon for peptide sequence deletion, \=post	AAAAQSRVVRV\LS\ AAAAQSRVVRV\LS\ AAAAQSRVVRV\LS\ AIATSVCHGPPCRQL IG\KGGRSS\VSGIVAT LRRYVVNH\LGRMG CDKYDIMHL\RPMGD EWDARDKDSIRRVV NL\IGTRP/WKSKNSD IPQAIAQLSK\EAG\V HLNANIKSSSRYLRN EEMKKDLLSKKEPE QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
MSRSAITA HHALMPH GFGA\TGF SQVIIPYR LGQLLFLE QHSNVVIN YEDVFVK EKFIHV\SH KNRKREK PEDLENSC TKSVAEQI YLGRSQE 15547 45915 A 15637 560 1394 RKSSSVHN RHLGRCR DLSAATS\ KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	AIATSVCHGPPCRQL IG\KGGRSS\VSGIVAT ICRRYVVNH\LGRMG CDKYDIMHL\RPMGD EWDARDKDSIRRVV NL\IGTRP/WKSKNSD IPQAIAQLSK\EAG\V HLNANIKSSSRYLRN EEMKKDLLSKKEPE QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
MSRSAITA HHALMPH GFGA\TGF SQVIIPYRI LGQLLFLE QHSNVVII YEDVFVK EKFIHV\SF KNRKREK PEDLENSQ TKSVAEQI YLGRSQE 15547 45915 A 15637 560 1394 RKSSSVHI RHLGRCRI DLSAATS\ KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	AIATSVCHGPPCRQL IG\KGGRSS\VSGIVAT ICRRYVVNH\LGRMG CDKYDIMHL\RPMGD EWDARDKDSIRRVV NL\IGTRP/WKSKNSD IPQAIAQLSK\EAG\V HLNANIKSSSRYLRN EEMKKDLLSKKEPE QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
HHALMPH GFGA\TGF SQVIIPYRG LGQLLFLE QHSNVVII YEDVFVK EKFIHV\SH KNRKREK PEDLENSG TKSVAEQI YLGRSQE YLGRSQE OLSAATS\ KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	IG\KGGRSS\VSGIVAT LRRYVVNH\LGRMG CDKYDIMHL\RPMGD EWDARDKDSIRRVV NL\IGTRP/WKSKNSD IPQAIAQLSK\EAG\V HLNANIKSSSRYLRN EEMKKDLLSKKEPE QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
GFGA/TGF SQVIIPYRG LGQLLFLE QHSNVVIN YEDVFVK EKFIHV\SH KNRKREK PEDLENSG TKSVAEQI YLGRSQE 15547 45915 A 15637 560 1394 RKSSSVHN RHLGRCRG DLSAATSV KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	CLRRYVVNH\LGRMG CDKYDIMHL\RPMGD EWDARDKDSIRRVV NL\IGTRP/WKSKNSD IPQAIAQLSK\EAG\V HLNANIKSSSRYLRN EEMKKDLLSKKEPE QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
SQVIIPYRE LGQLLFLE QHSNVVII YEDVFVK EKFIHV\SH KNRKREK PEDLENSQ TKSVAEQI YLGRSQE 15547 45915 A 15637 560 1394 RKSSSVHN RHLGRCRE DLSAATSY KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	CDKYDIMHL\RPMGD EWDARDKDSIRRVV NL\IGTRP/WKSKNSD IPQAIAQLSK\EAG\V HLNANIKSSSRYLRN EEMKKDLLSKKEPE QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
LGQLLFLE QHSNVVIN YEDVFVK EKFIHV\Si KNRKREK PEDLENSQ TKSVAEQI YLGRSQE 15547 45915 A 15637 560 1394 RKSSSVHN RHLGRCRI DLSAATSY KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	EWDARDKDSIRRVV NL\IGTRP/WKSKNSD IPQAIAQLSK\EAG\V HLNANIKSSSRYLRN EEMKKDLLSKKEPE QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
QHSNVVIN YEDVFVK EKFIHV\SH KNRKREK PEDLENSQ TKSVAEQI YLGRSQE YLGRSQE OLSAATSY KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	NL\IGTRP/WKSKNSD IPQAIAQLSK\EAG\V HLNANIKSSSRYLRN EEMKKDLLSKKEPE QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
YEDVFVK EKFIHV\SH KNRKREK PEDLENSQ TKSVAEQI YLGRSQE 15547 45915 A 15637 560 1394 RKSSSVHN RHLGRCR DLSAATSY KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	IPQAIAQLSK\EAG\V HLNANIKSSSRYLRN EEMKKDLLSKKEPE QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
EKFIHV\SI KNRKREK PEDLENSQ TKSVAEQI YLGRSQE 15547 45915 A 15637 560 1394 RKSSSVHN RHLGRCRI DLSAATSY KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	HLNANIKSSSRYLRN EEMKKDLLSKKEPE QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
KNRKREK PEDLENSC TKSVAEQI YLGRSQE 15547 45915 A 15637 560 1394 RKSSSVHN RHLGRCRI DLSAATSY KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	EEMKKDLLSKKEPE QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
PEDLENSC TKSVAEQI YLGRSQE 15547 45915 A 15637 560 1394 RKSSSVHN RHLGRCRG DLSAATSY KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	QLIHIAKNEKACFEEN PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
TKSVAEQI YLGRSQE 15547 45915 A 15637 560 1394 RKSSSVHN RHLGRCR DLSAATSY KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	PFDEEIVDATHQLTS NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
15547 45915 A 15637 560 1394 RKSSSVHN RHLGRCRI DLSAATSY KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	NAASGLLGSRAPGAH CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
15547 45915 A 15637 560 1394 RKSSSVHN RHLGRCRI DLSAATSV KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
RHLGRCRO DLSAATSY KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	CRHRLVSPSPAMAE YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
DLSAATSY KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	YTEDDFYCPVCQEVL ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
KTPVRTTA MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	ACQHVFCRKCFLTA CPLCRGNVTRRERA LENIMR\KFSGSCRCC
MRESGAH CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	CPLCRGNVTRRERA LENIMR\KFSGSCRCC
CPERALDI AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	LENIMR\KFSGSCRCC
AK\QIKFY DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	
DEYGVSSI NNTGIKQI QRNAKTE GSGPNLGI	RMRHHYK\SCKKYQ
NNTGIKQI QRNAKTE GSGPNLGI	IPNFQISQDSVGNRIE
QRNAKTE GSGPNLGI	PGAPAFGDGRVGCR
GSGPNLGI	LLAAAGESPCTMSL
	PSPRSKSSSVTSPQPC
	FVSFPVHYVLEGSR
	LKTDRIFGH
	EVKFEKQCPEAVLLH
1 1 1 1 1	DENKRCDLREPRKPE
	LLVPTTQPGRNEGQS
1 1 1 1	RCSKTSCCLSGKEPS
1 1 1 1 1 1	.VPVQNTTIKPDSGIR
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	ETFTMSTALDVLQM
1 1 1 1	FLAAGTHLGGTDLD
1 1 1 1 1	CKRKSGGIYIINLKRS
1 1 1 1 1 1 1	APAIVAIENPADVSV
ISSSSTGQI	RAMLKCAAATGATP
1 1 1 1 1 1 1 1	TFTNQIQATFPNPRV
1 1 1 1	TGHQPLAEASYV/NN
	TDSPLHHVDIAIPCN
R/KAHSVD	LMWMLAREILPM\R
GTIARKHY	WWE\DPEEIEKEEQA
AADKALT	KEEFQGE
	NKGAHSVG*MWR
	RMRGTISGEHPWEV
1 1 1 1 1	RDTEEIEKEEQAAAD
	FQGEWTAPAPEFTAT
1 1 1 1	SEGVQVPSVPIQQFP
	PATEDWSAAPTAQA
TEWVGAT	
15550 45918 B 15640 88 531	
15551 45919 A 15641 1 1302	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15552	45920	A	15642	150	957	KAGASCCGSANPYVSVGKSCV LLAMAQLQTRFYTDNKKYAVD DVPLSIPAASEIADLSNIINKLLK DKNEFHKHVEFDFLIKGQFLRM PLDRHMEMENISSEEVVEIEYV EKYTAPQARSNAWFHDDWISSI KRGRG\RILTGSYDKTSRIWSLE GKSIMTIVGHTDVVKDVAWVK KDSLSCLLLSASMDQTILLWEW NVERNKVKALHCCKGHAGSV NSIAVDGSGTKYFGLRIEPKPA VSLVEEINSLSYRWTADNYKV GSGGKEEED
15553 15554	45921 45922	A	15643	335	537	VFCCPHCSLTFSKSFSSFSRLVR HQQTHWKQKSYLCPICDLSFGE KEGLMD/HLEGL*RQGPGQSSH HKCRVILGQWLGFSHDVPTMA GEEWKHGGDQSPPRIHTPRRRG LREKACKGDKTKEAVSILKHK VYFKQVTLDSSCRL*ERQVKST
						SSWNILSSTWRLPFCPTSVVLRF LTWICLRTISTSSISWFPADSS
15555	45923	Α	15645	153	828	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15556	45924	A	15646	1	2002	MLESAGNQDIEDVDIVLRQIQV
						KNRSTTEVGQNGNLQVEVSYT
	1					QAHIAPRPLASQGTWPHFVSSA
						PQPCFQVQNHSKYQNLYYSGFI
						RETEPISKGDVGEWLSAGKGES
						SAMFASEQEISKDEQGTPVLGS
						FYWEVDSPRKESSQAWAPGQE
						WIKLERDTTEEKMFEQLKPIEP
						VQKTLPWVGEVAATLQEAMK
						RDCWREARVKKKPVTFEDVAV
						NFTQEEWDCLDASQRVLYQDV
						MSETFKNLTSVEGKKKELREQ
						HPSLRDEGTSDDKVFLACRGA
						GQCPLSAPAGTMDRTRVLQAS
ļ						QAGPP/FFFATTVGKCFSRRSYL
						YSHQFVHNPKLTNSCSQCGKLF
						RSPKSLSYHRRMHLGERPFCCT
						LCDKTYCDASGLSRHRRVHLG
l						YRPHSCSVCGKSFRDQSELKRH
				}		QKIHQNQEPVDGNQECTLRIPG
		Ī				TQAEFQTPIARSQRSIQGLLDVN
						HAPVARSQEPIFRTEGPMAQNQ
						ASVLKNQAPVTRTQAPITGTLC
ļ						QDARSNSHPVKPSRLNVFCCPH
						CSLTFSKKSYLSRHQKAHLTEP
1						PNYCFHCSKSFSSFSRLVRHQQ
						THWKQKSYLCPICDLSFGEKEG
İ						LMDHWRGYKGKDLCQSSHHK
						CRVILGQWLGFSHDVPTMAGE
				:		EWKHGGDQSPPRIHTPRRRGLR
				•		EKACKGDKTKEAVSILKHK
15557	45925	Α	15647	1	1048	MPSLAELFEGSSKAFGAAASQL
						NFPVASGEMLKFKYG\LRNPLH
						AVAAEPIASRTSRLNLFFQGKPP
						FMTQQQMSPLSREGILDALFVL
						FEECSQPALMKIKHVSNFVRKY
						SDTIAELQELQPSAKDFEVRSLV
						GCGHFAEVQVVREKATGDIYA
						MKVMKKKALLAQEQVSFFEEE
1						RNILSRSTSPWIPQLQYAFQDK
						NHLYLVMEYQPGGDLLSLLNR
						YEDQLDENLIQFYLAELILAVH
						SVHLMGYVHRDIKPENILVDRT
						GHIKLVDFGSAAKMNSNKMVN
						AKLPIGTPDYMAPEVLTVMNG
						DGKGTYGLDCDWWSVGVIAY
			<u> </u>	<u> </u>	L	EMIYGRSPFAEGTSARTFNNIM

NO: of peptide sequence hod in USSN location of first codon for last amino acid to sequence logically codon for peptide of peptide sequence do	Amino acid sequence (X=Unknown, '=Stop codon, /=possible nucleotide leletion, \=possible nucleotide insertion)
sequence	referion, (-possible nucleotide insertion)
	KRQQAFGLLDLHQWFARGSQG
	FRLQTEGCIIGFPTFEV*RHSGFL
	SCQSSCTGSFSSVWADVPLMFE
	GAAQPV*PVPSRGSADFRPLSIC
	RSSRILLVKRCENSGVR
	TMKSQRQIIKKAVTTKEMKSV
	CRRD/TPAFMFIAALFKIAKIRIQ
	RLCLSTDEWIKKMWYIHTMEY
	YSTLK
15560 45928 B 15650 161 3156	
15561 45929 A 15651 14 270	
15562 45930 B 15652 787 1821	
	MAKSKGCSAATLASTHQILVTP
	WSCDNQKCLQTLPNVPGGTKL
	LCEDHWFRNDSCNQAIPTIAT
	LLSFPSLQASLWTTHFLSPLNH
	HPVNPDSHLEQPPGQSPRLNFE
	AGDPERAPHALGCKRSRTRATS
	DRQPMASRDPGWGKASVGKA
	CALGSAVLPAGNDCTTDNLGL
	GSAGFEDLRLRLGQGPPRRQGA
	LQAAPRKASSCAAGQRLAPDSS
	FSRKRRPSRQRLGCETAAVADT
	AEAKSSEAPAFGWSTMDEAGS
	SASGGGFRRALDEPPNSRIFLVI
	SKYTPESVLRERFSPFGDIQDIW
1 1 1 1 1 1	VVRDKHTKESKGIAFVKFARSS
	QACRAMEEMHGQCLGPNDTKP
	KVRVPGSGCPRGKEWASWTSL
	HLLLCRGEGGVEHPFPAADQT
	APGLGTGAALRRGSPPTKLRRV
	AGKGTGCGVFAPRERTRQPRPP
	GDRTAVAGLP\SPSPSLIHQLLSS
	LPNSS*PTLSTFLTFLSQMPRFP
	YIKNGKTL/PFPQVFIAQSRSSGS
	HRDVEDEELTRIFVMIPKSYTEE
	DLREKFKVYGDIEYCSIIKNKVT
	GESKGLGYVRYLKPSQAAQAIE
	NCDRSFRAILAEPKNKASESSE
	QDYYSNMRQEALGHEPRVNMF
	PFGHGVVQYFNVASAIYAKYK
	LHGFQYPPGNRIGVSFIDDGSN
	ATEKMATQMVAAQLASMVWN
	NPSQQQFMQFGGSSGSQLPQIQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ļ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15564	45932	A	15654	375	2059	DSSVWQTPQKQRAVRPLHSGG
1						APWTKLAALRATRGFRPGVDS
						LDEPPNSRIFLVISKYTPESVLRE
ļ						RFSPFGDIQDIWVVRDKHTKES
						KGIAFVKFARSSQACRAMEEM
						HGQCLGPNDTKPIKVRVPGSGC
						PRGKEWASWTSLHLLLCRGEG
						GVEHPFPAADQTAPGLGTGAA
						LRRGSPPTKLRRVAGKGTGCG
						VFAPRERTRQPRPPGDRTAVAG
						LPLSLALTNPPALEFTAKLVLA
						YFEYFSDLPLSDAQISIYKKRKN
						STFPQVFIAQSRSSGSHRDVEDE
		ŀ				ELTRIFVMIPKSYTEEDLREKFK
						VYGDIEYCSIIKNKVTGESKGL
		1				GYVRYLKPSQAAQAIENCDRSF
						RAILAEPKNKASESSEQDYYSN
						MRQEALGHEPRVNMFPFGHGV
						VQYFNVASAIYAKYKLHGFQY
						PPGNRIGVSFIDDGSNATDLLIK
		ŀ				MATQMVAAQ/RLHQWCGITQV
						SNNL/WQFGGSSGSQLPQIQTD
						VVLPSCKKKAPAETPVKERLFI
		ļ.				VFNPHPLPLDVLEDIFCRFGNLI
						EVYLVSGKNVGYAKYADRISA
		ļ				NDAIATLHWKDSEWGET
15565	45933	A	15655	257	412	HA A ODGGCVECK GODY GYAN DA
15566	45934	Α	15656	3	866	HAASPCCCYEEKQSPLQKHLRV
						CSLRTWSLAARVAEGTAETVD
						PSAAPKTSVWSSRVCWGRLLA
						TISPSVNETDTCGMEDRRLVVC
		į				RPWRAQCHMKAWGWRHPEEG
						GTHSGAPSAVLQALAVAIQLGG
		ł				HLADPLLQVDPLSSCGAVSLDI
						LIYLVFYYRTASVPETYIVKTLF
						KKLESQSLIQSNVLTRSNSMKA ERGEEAAKEKSEASRGWFMRF
						KERSCLHNIKMQGEAASADVE
						AAASYPKDLAKITDEHGYTKQ Q\FDVYKTAFCWRVMLKKYVK
1						1 -
L				<u> </u>	l	LLFVLCPLP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
15567	45935	A	15657	333	1034	RLSTTVMRGAFVF\MHKMPRL WLDYCQF\LMDQGRVTHTRRT FDRALRALPITQHSRIWPLYLRF LRSHPLPETAVRGYRRFLKLSP ESAEEYIEYLKSSDRLDEAAQR LATVVNDERFVSKAGKSNYQL WHELCDLISQNPDKVQSLNVD AIIRGGLTRFTDQLGKLWCSLA DYYIRSGHFEKIRKAPVNGGPT WMVGQNSVPRFQTRRLTPLKR KGPLVTDFREKQPTPLAQE
15568	45936	A	15658	172	346	ARPCRWCPGS*TSASAPGPGLR PCPP\GGDRTPASPPLHNQAAA APPGPGTWRGAAAS
15569	45937	Α	15659	293	482	
15570	45938	A	15660	468	791	LAKCGCLEARGRARQTSGRPRP TDGGP*SPCWETGKGHPVLGR RGS/PGKQGL*SLPQSCQGQWR CAASRLLSSKNSWACMGSREA VSQETGTSSSSLGLMRPSDKKP
15571	45939	Α	15661	366	609	
15572	45940	В	15662	321	739	
15573	45941	Α	15663	1	1815	
15574	45942	Α	15664	i	1911	
15575	45943	A	15665	437	1125	PVSCSGHSRWAAAAISIALSMIE TVNMVFTFYSRHHGTISTRVL/I GKRPERQDGQDTVNLGTLEET ASPSPVVTTSPPQPMLPSAFPPL SEEINPALPEATVMASPEAVAR QDNVDSAQEPPPTPLCASRPTT RQKSWWGARGEVESVTHEEM HYTGKELLEFSNLYKEKSGEQA WEWILRVWDNGGRNIQLDHPE FVDLGPLSRNSAFNVAVRGVK KGSKSLFAWLGEI
15576	45944	С	15666	49	186	
15577	45945	A	15668	80	1218	GAERLPEGPGKRMVFRRFVE\V GRVAYVSFGPHAGKI/VSAMVD VF\DQNKGFWSNGPCTQVR\RQ A\MPFK\CMQLT\DFILKFPHSAH Q\KYVRQSLGRREDINTKLGQP TRWGQGRFESQEEKEKPRLTDF *SVF*S**RAKEN*RDRINPRNE V*EAFKRAAFPGKASSQKKAPG Y*GVLLAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
15578	45946	A	15668	253		
15579	45947	Α	15669	1	1819	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ļ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
15580	45948	Α	15670	1	2422	MCPILREHRGSTEGKRRQKRLP
						LELTLKDEEVLAHRQEIEVAKA
						SPSKGNIIIHFGYKGVEYHDQQ
						DVTSNFLGAMWLISITFLSIGYG
ļ						DMVPNTYCGKGVCLLTGIMGA
						GCTALVVAVVARKLELTKAEK
			!			HVHNFMMDTQLTKRHCPWAA
						TIQPKLSFLISLPGCDYRGESAR
						ARPLSHPPLQLGLGKDARYLRI
						CDRSLLCVCAAAGNAGLRPRD
						SPHTLLHWRSRSPYFFSPKPVPP
						QLLGGQPAAHSRRDDLGEAGV
						GRGGCCFAALRGVQSEKWSGF
		ŀ				HTVPEPAPGQGDVMPIVLVRPT
						NRTRRLDSTGAGMGPSSHQQQ
						ESPLPTITHCAGCTTAWSPCSFN
						SPDMETPLQFQRGFFPEQPPPPP
						RSSHLHCQQQQQSQDKPCPPFA
						PLPHPHHHPHLAHQQPASGGSS
						PCLRCNSCASSGAPAAGAGDNL
						SLLLRTSSPGGAFRTRTSSPLSG
						SSCCCCCSSRRGSQLNVSELTP
						SSHASALRQQYAQQSAQQSAS
1						ASQYHQCHSLQPAASPTGSLGS
						LGSGPPLSHHHHHPHPAHHQH
<u> </u>						HQPQARRESNPFTEIAMSSCRY
						NGGVMRPLSNLSASRRNLHEM
						DSEAQPLQPPASVGGGGGASSP
						SAAAAAAAVSSSAPEIVVSKP
						EHNNSNNLALYGTGGGGSTGG
						GGGGGGSGHGSSSGTKSSKKK
		<u> </u>				NQNIGYKLGHRRALFEKRKRLS
15581	45949	Α	15671	1	246	SEPPHRGPTGALPSGAVRRGPL
						SSRPQIGRSTDSLHCVPGKATD
						AQRQPVKAAGSGAVPCKATG
15500	45050	_	15.650	12	741	WSCPKL*EPISCINVTIC
15582	45950	В	15672	43	741	ELIZODDENAMORI ODEOLOMANA
15583	45951	Α	15673	687	1012	FHTSRPFWNISELSRFCIQKYKK
						PWHSVSDFYRYQFFICQGTACC
						PASLRM*HCPN*ILQGTSSQNCC
						SCSGKSCSPMQQRHRSTVCSGS
		<u> </u>				GTFCVGSCESFEGTYFSR